

1 **A Geospatiotemporal and Causal Inference Epidemiological Exploration of Substance**
2 **and Cannabinoid Exposure as Drivers of Rising US Pediatric Cancer Rates**

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4
5 *Short Title:*

6 *Geospatiotemporal Cannabinoid Exposure and Total Pediatric Cancer Incidence*

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27 Word Count: 4,592.

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30 Key words: cannabis, cannabinoid, Δ^9 -tetrahydrocannabinol, cannabigerol, genotoxicity,
31 acute leukaemia, pediatric cancer

32

33 Abstract

34

35 Background. Age-adjusted US total pediatric cancer incidence rates (TPCIR) rose 49% 1975-
36 2015 for unknown reasons. Prenatal cannabis exposure has been linked with several pediatric
37 cancers which together comprise the majority of pediatric cancer types. We investigated
38 whether cannabis use was related spatiotemporally and causally to TPCIR.

39

40 Methods. State-based age-adjusted TPCIR data was taken from the CDC Surveillance,
41 Epidemiology and End Results cancer database 2003-2017. Drug exposure was taken from
42 the nationally-representative National Survey of Drug Use and Health, response rate 74.1%.
43 Drugs included were: tobacco, alcohol, cannabis, opioid analgesics and cocaine. This was
44 supplemented by cannabinoid concentration data from the Drug Enforcement Agency and
45 ethnicity and median household income data from US Census.

46

47 Results. TPCIR rose while all drug use nationally fell, except for cannabis which rose.
48 TPCIR in the highest cannabis use quintile was greater than in the lowest (β -estimate=1.31
49 (95%C.I. 0.82, 1.80), $P=1.80 \times 10^{-7}$) and the time:highest two quintiles interaction was
50 significant (β -estimate=0.1395 (0.82, 1.80), $P=1.00 \times 10^{-14}$). In robust inverse probability
51 weighted additive regression models cannabis was independently associated with TPCIR (β -
52 estimate=9.55 (3.95, 15.15), $P=0.0016$). In interactive geospatiotemporal models including
53 all drug, ethnic and income variables cannabis use was independently significant (β -
54 estimate=45.67 (18.77, 72.56), $P=0.0009$). In geospatial models temporally lagged to 1,2,4
55 and 6 years interactive terms including cannabis were significant. Cannabis interactive terms
56 at one and two degrees of spatial lagging were significant (from β -estimate=3954.04
57 (1565.01, 6343.09), $P=0.0012$). The interaction between the cannabinoids THC and
58 cannabigerol was significant at zero, 2 and 6 years lag (from β -estimate=46.22 (30.06, 62.38),
59 $P=2.10 \times 10^{-8}$). Cannabis legalization was associated with higher TPCIR (β -estimate=1.51
60 (0.68, 2.35), $P=0.0004$) and cannabis-liberal regimes were associated with higher
61 time:TPCIR interaction (β -estimate= 1.87×10^{-4} , (2.9×10^{-5} , 2.45×10^{-4}), $P=0.0208$). 33/56
62 minimum e-Values were >5 and 6 were infinite.

63

64 Conclusion. Data confirm a close relationship across space and lagged time between
65 cannabis and TPCIR which was robust to adjustment, supported by inverse probability

66 weighting procedures and accompanied by high e-Values making confounding unlikely and
67 establishing the causal relationship. Cannabis-liberal jurisdictions were associated with
68 higher rates of TPCIR and a faster rate of TPCIR increase. Data inform the broader general
69 consideration of cannabinoid-induced genotoxicity.

70

71 Introduction

72

73 CDC Surveillance, Epidemiology and End Results (SEER) data from 9 US cancer registries
74 indicates that the age-adjusted total Pediatric (age less than twenty years) cancer incidence
75 rate (TPCIR) has risen 49.0% from 12.96 to 19.32 / 100,000 from 1975-2015 [1]. Cancer
76 incidence is U-shaped across the pediatric age range being higher in the under 5 years and
77 over 14 years age groups [2]. Leukaemias, brain and nervous system, neuroblastoma, soft
78 tissue sarcoma, lymphoma and testicular cancer are amongst the commonest pediatric cancers
79 [2, 3].

80

81 Notwithstanding a generally falling mortality rate from childhood cancer, the TPCIR
82 incidence is acknowledged to be rising since the records of collated cancer registries were
83 first published in 1975 [2]. The cause of this unprecedented increase is at present unclear.
84 Moreover major ethnic differentials are evident for tumours such as All Childhood Cancer
85 (ACC), acute lymphatic leukaemia (ALL) and brain and testicular cancers where the rates in
86 African-American patients vary from 20-70% of those in the Caucasian-American
87 community [2]. Again the reasons for such large ethnic disparities are unknown. It therefore
88 appears that several of the major questions relating to the aetiopathogenesis of pediatric
89 cancer are outstanding.

90

91 Whilst in adult populations the relationship between cannabis use and cancer incidence is
92 controversial with both positive and negative reports in existence [4, 5], amongst pediatric
93 populations the situation is much clearer. It was noted by the California Environmental
94 Protection Agency in a very detailed literature review that five of six studies reported a
95 positive relationship [6-11]. Parental cannabis use has been linked with acute lymphatic
96 leukaemia, acute myeloid leukaemia, childhood astrocytoma, rhabdomyosarcoma and
97 neuroblastoma [2, 7-12]. Together these comprise 60-70% of the total cancers seen in
98 children younger than 14 years and those between 15 and 20 years [2]. In such a context it
99 becomes plausible that the rise in cannabis use since the 1960's may be a primary driver of
100 total pediatric cancer.

101

102 Testicular cancer is a particularly interesting case. It is well established that testicular cancer
103 occurs mainly in younger men with an age peak at 30-34 years and 20% of cases occur in the
104 pediatric age range [1]. The testes houses the germ cells and cannabinoids are known to have

105 myriad direct effects on the reproductive tract in both sexes [13-17]. There is great
106 uniformity in studies of the cannabis-testicular cancer link as all four studies found a risk
107 elevation of over two-fold [18-21] with an overall risk for current, weekly and chronic
108 smokers of non-seminomatous germ cell tumours estimated in meta-analysis of 2.59
109 (95%C.I. 1.60-4.19) [22]. Since pediatric cancer often results from inherited genetic errors
110 [23, 24] this implies that major genetic errors in germ cells are induced by parental cannabis
111 exposure.

112

113 Adding to concerns related to the potentially genotoxic actions of prenatal cannabinoid
114 exposure (PCE) is an increasing interest in elevation of many birth defects following PCE in
115 Hawaii, Colorado, Canada and Australia [25-28]. A recent report noted a three-fold rise in
116 total congenital defects in the northern Territories of Canada where more cannabis is smoked
117 [28]. Down's syndrome, due to a major genetic trisomic error, has also been found to be
118 elevated following PCE in Hawaii, Colorado and Australia [25-27] and this syndrome has an
119 established link with childhood ALL with 6-10% of Down's syndrome children being affected
120 by this malignancy [29, 30].

121

122

123 As discussed below the physiology and pathophysiology of both the endocannabinoid system
124 and the impacts of diverse exogenous phytocannabinoids is presently being studied in great
125 detail and major impacts on reproductive health, genetic and physiological quality of
126 gametes, epigenetic effects on both DNA methylation and histone synthesis and signalling,
127 immunomodulatory and mitochondriopathic effects, and transgenerational inheritable
128 epigenetic effects in both man and mouse are well established and have been demonstrated by
129 a number of investigators [15, 17, 31-38].

130

131 Concerns are heightened by the recent demonstration that 69% of cannabis dispensaries in
132 Colorado recommended cannabis use to pregnant patients for various symptoms in a recent
133 telephone survey [39] and that in 2017 an estimated 161,000 women used cannabis whilst
134 pregnant across USA [40, 41].

135

136 Taken together these data suggest that an improved understanding of cannabis-related
137 carcinogenesis in the closely defined pediatric context might well lead to important insights
138 into cannabis-related genotoxicity more generally [42, 43]. Moreover the advent of

139 sophisticated geospatial analysis together with some of the formal techniques of causal
140 inference analysis implies that sophisticated and modern analytical procedures could be
141 brought to bear on these important and increasingly topical issues. Techniques such as
142 inverse probability weighting and e-Values are designed to formally investigate causal, as
143 opposed to merely associational, relationships.

144

145 The objective of this study was to determine if the rise in pediatric cancers across USA
146 paralleled the recent rise in the use of cannabis when considered formally across space and
147 time, and if the relationship met the criteria for causal inference when assessed by strict
148 quantitative criteria.

149

150

151 Methods

152

153 Data. Annual data on age-adjusted rates of pediatric cancer cases occurring in patients less
154 than 20 years old was accessed from the publicly available SEER*Explorer website [1]. Data
155 on state-based pediatric cancer rates was accessed via the SEER*Stat software from the
156 SEER / NCI database [44]. Drug use data was accessed from the nationally representative
157 National Survey of Drug Use and Health (NSDUH) conducted by the Substance Abuse and
158 Mental Health Services Administration (SAMHSA) [45]. This survey reports a 74.1%
159 response rate [46]. Data on the following drug variables was collated: monthly cigarette use;
160 annual alcohol use disorder, monthly cannabis use, annual analgesic abuse and annual
161 cocaine use. Data on ethnic composition and median household income by state and year
162 was accessed via the tidycensus package in R from the US Census Bureau. The ethnicities
163 for which data was collected were: Caucasian American, African American, Hispanic
164 American, Asian American, American Indian / Alaskan Native American, Native Hawaiian /
165 Pacific Islander American. Data on national cannabinoid concentrations for Δ^9 -
166 tetrahydrocannabinol (THC), cannabiol, cannabigerol and cannabichromene was obtained
167 from various published reports [47-49]. Data on cannabis legal status was adduced from an
168 internet search [50].

169

170 Derived Data. Given the clear differences in drug use by ethnicity it was considered
171 important to formally take ethnic cannabis use into account in regression modelling. Data on
172 the frequency of cannabis use by ethnicity was available at the national level from the
173 SAMHSA Substance Abuse and Mental Health Data Archive (SAMHDA) Restricted Use
174 Data Analysis System (RDAS) [45]. For each ethnicity and for each year the percentage of
175 the ethnicity using cannabis at the midpoint of the indicated frequency were multiplied
176 together and summed to gain an ethnic cannabis use index. Hence if fraction x of an ethnicity
177 used cannabis from 20-30 days per month then x would be multiplied by 25. This was
178 repeated and summed across all use frequencies to obtain a specific ethnic cannabis use index
179 for that year. This index was multiplied by the state cannabis use rate and the THC
180 concentration in that year to derive an estimate of the ethnic exposure to THC in each state.
181 Similarly the concentration of selected cannabinoids was multiplied by the state cannabis use
182 rate to derive a state based exposure to that cannabinoid. Cannabis use quintiles were defined
183 in each year and concatenated to form strata across all years.

184

185 Missing data. The total pediatric cancer rate for Wyoming 2008 was absent. This was
186 imputed as the mean of its rate in 2007 and 2009. The rate of analgesic use was missing for
187 all states in 2015. This was imputed as the mean of the state rates for 2014 and 2016.

188

189 Statistics. R version 4.0.2 (2020-06-22) from CRAN was used for data analysis and accessed
190 via the RStudio 1.2.5042 (2009-2020) GUI. Data analysis was performed in September
191 2020. Graphs and map-graphs were drawn using packages ggplot, albersusa and sf.

192 Covariates were log-transformed to approximate normality based on the Shapiro-Wilks test.

193 Linear, mixed effects, panel, robust marginal structural models and spatial models were
194 studied using packages base, nlme, plm, survey and splm (spatial panel linear models)

195 respectively [51-53]. In each case model reduction was performed by the classical technique

196 of serial deletion of the least significant term. A variety of modelling procedures was

197 employed for the following reasons. Mixed effects regression was useful for state-wise study

198 of data, for inverse probability weighted corrections, and for generation of standard

199 deviations which can be input to eValue calculations. Panel regression modelling was well

200 suited to the time series sequential nature of the dataset, can be inverse probability weighted

201 and allowed the use of both lagging and instrumental variables. Robust regression was

202 conducted to examine the robust effects after inverse probability weighting. Spatiotemporal

203 regression was performed as the data are inherently distributed across space and time and

204 there was good evidence from the models for both spatial and temporal autocorrelation (see

205 Results). As the models also produce a variance estimate their output is well suited to the

206 calculation of e-Values. Inverse probability weighting was conducted with the ipw package

207 and e-Values for regression models were calculated with the package EValue. Tests for trend

208 were conducted with the chi squared test in Base. T-tests were conducted for parametric

209 group comparisons and were two tailed. $P < 0.05$ was considered significant throughout.

210

211 Panel analysis utilized the pooling technique, a time effect, the random method of Swamy,

212 the instrumental method of Amemiya and were inverse probability weighted. Robust

213 structural models were conducted by state and were inverse probability weighted.

214

215 Spatial analysis. Interstate geospatial linkages were made on the “queen” basis of shared

216 edges or corners and compiled with the poly2nb function from package spdep. They were

217 edited as described so that no state, such as Alaska or Hawaii, was left geospatially isolated

218 (as shown in Results). Model specification of spatial models was undertaken from the

219 general full model to the specific [54]. That is to say the standard spatiotemporal regression
220 model was conducted using the `splm` function `spreml` (spatial panel random effects maximum
221 likelihood) including spatial autocorrelation after Kapoor, Kelejian and Prucha [55], random
222 effects, serial correlation in the residual errors and spatial autocorrelation, coded as `sem2srre`
223 in `spreml` models [52]. Significance of the final model parameters ϕ , ψ , ρ and λ
224 which quantify random error, serial correlation in the residuals, spatial error correlation and
225 spatial autocorrelation respectively, confirmed that this maximal structure was appropriate
226 (see Results tables). The spatial error adjustment of Kapoor, Kelejian and Prucha takes into
227 account spatial correlation in both the exposure and the outcome and this was considered to
228 be reflective of the real world situation in this case [54]. `spreml` models do allow the use of
229 both spatial and temporal lagging which has been utilized as described. At the time of
230 writing `splm` and `spreml` spatial models do not allow the use of instrumental variables or
231 inverse probability weighting which implies the need for supplementary techniques.

232

233

234 Causal inference. Two techniques of causal inference were employed. Inverse probability
235 weights were constructed for the exposure of interest, monthly cannabis exposure, as a
236 function of the other drug variables which were our primary variables of interest. These
237 weights were used to weight mixed effects, panel and robust regression models appropriately.
238 The effect of this procedure is to equalize exposure across study groups and has also been
239 validated for continuous exposures as considered here. Such techniques are said to create
240 pseudo-randomized groups from which causal inferences can properly be made. We also
241 calculated e-Values which are a measure of the association required of any unmeasured
242 potential confounder variable with both the exposure and the outcome to discount the
243 reported results. In the literature minimum (of the two) e-Values above 1.25 are commonly
244 considered of relevance [56].

245

246

247 Data availability. All data, including R code, inverse probability weights, geospatial weights,
248 and source datasets, has been made publicly available through the Mendeley data base
249 repository and may be accessed at this URL: <http://dx.doi.org/10.17632/cnww9hdspd.1> .

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251

252

253 Ethics. The datasets used were all publicly available and de-identified. No reference has
254 been made at any point to individually identifiable data. The present work was approved by
255 the University of Western Australia Human Research Ethics Committee on June 7th 2019
256 (No. RA/4/20/4724).

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263 Results

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265 Inspection of the SEER*Explorer website shows that at the national level that age-adjusted
266 rates of several cancers in the pediatric age group (younger than 20 years) are rising including
267 all cancer and acute lymphatic leukaemia which is the commonest tumour. The annotation
268 on the SEER website is made from the JoinPoint program which also comes from NCI and
269 CDC. These tumours are listed in Table 1 and illustrated in Figure 1 using data based on 9
270 US cancer registries 1975-2017. Supplementary Figure 1 shows other cancers which are
271 mostly rising utilizing data from 21 US cancer registries 2000-2017.

272

273 Figure 2 shows national drug exposure data from NSDUH 2003-2017 and US Census bureau
274 median household income data. It is important to note that exposure to most classes of drugs
275 is dropping with the notable exception of cannabis. Since SAMHSA NSDUH data could be
276 temporally matched to the CDC SEER cancer database for the years 2003-2017, this became
277 the period of analysis.

278

279 Figure 3 shows the concentration of various cannabinoids found in federal cannabis seizures
280 1980-2017 [47-49].

281

282 Figure 4 shows the age-adjusted state-based TPCIR plotted as a function of exposure to the
283 various substances listed. The regression line for cannabis is noted to be weakly and non-
284 significantly positive.

285

286 Figure 5 shows plots of the TPCIR rate against selected cannabinoids. The regression lines
287 for THC and cannabigerol appear to be strongly positive.

288

289 Figure 6 shows the TPCIR as a function of ethnic cannabis exposure. In each case the
290 regression line appears to be strongly positive and up-sloping.

291

292 Table 2 lists applicable results from linear regression against time, cannabis, THC, various
293 substances, cannabinoids and ethnicity. Many results are significant with the notable
294 exception of cannabis.

295

296 Figure 7 shows the result of assessing the TPCIR as a function of cannabis use quintiles both
 297 cross-sectionally (boxplots) and over time (scatterplots). Panel A appears to show a rising
 298 trend with cannabis use quintile. One notes in particular that the notches of the fourth and
 299 fifth quintiles do not overlap those of Quintiles 1 and 2 which indicates significance. In
 300 Panel B the highest two quintiles seem to be above the lower ones over time. Panel C and D
 301 look at the data dichotomized into the two highest quintiles compared to the three lower ones.
 302 Again in Panel C it is clear that the notches of the upper quintiles do not overlap those of the
 303 lower ones. Panel D shows that this holds true over time. Raw mean quintile data with
 304 standard errors appears in Supplementary Table 1.

305

306 When comparing the highest and lowest quintile of cannabis use the TPCIR in the highest
 307 quintiles is significantly greater than that in the lowest quintile ($t=5.038$, $df=299.6$,
 308 $P=8.15 \times 10^{-7}$). Comparing the two dichotomized cannabis quintile groups they are also
 309 significantly different ($t=5.641$, $df=673.6$, $P=2.4810^{-8}$). The chi squared test for trend across
 310 the quintiles does not reach significance ($\text{Chi.Squ.}=465.4$, $df=420$, $P=0.0623$). When these
 311 data are examined by linear regression the significant results shown in Table 3 are found.

312

313 **Table 3.: Linear Regressions on Quintiles**

314

315

Parameter Estimates			Model Parameters			
Parameter	Estimate (C.I.)	P-Value	R-Squared	F	dF	P-Value
<i>Quintiles</i>						
<i>lm(Cancer Rate ~ Quintile)</i>						
Quintile 2	0.2 (-0.29, 0.69)	0.4242	0.04527	9.34	4,745	2.27E-07
Quintile 3	0.14 (-0.35, 0.63)	0.5655				
Quintile 4	0.72 (0.23, 1.2)	0.0042				
Quintile 5	1.31 (0.82, 1.8)	1.8E-07				
<i>Dichotomized Quintiles</i>						
<i>lm(Cancer Rate ~ Dichotomized Quintiles)</i>						
Upper 2 Quintiles	0.9 (0.58, 1.22)	3.9E-08	0.0383	30.9	1,748	3.86E-08
<i>Dichotomized Quintiles Over Time</i>						
<i>lm(Cancer Rate ~ Year + Dichotomized Quintiles)</i>						
Upper 2 Quintiles	0.9 (0.59, 1.2)	1.1E-08	0.111	47.8	2,747	<2E-16
<i>lm(Cancer Rate ~ Year : Dichotomized Quintiles)</i>						
Lower 3 Quintiles	0.139 (0.1, 0.17)	1.2E-14	0.111	47.7	2,747	<2E-16
Upper 2 Quintiles	0.1395 (0.1, 0.17)	1.0E-14				

316

317 Table 4 presents results from increasingly complex robust inverse probability weighted
318 marginal structural models. Results for additive, interactive with drugs only, interactive
319 including drugs, race and income and interactive including cannabinoids, drugs, race and
320 income models are shown. It is particularly noteworthy that in a simple additive robust
321 model (listed first in the table) cannabis is independently highly significant (β -estimate=9.55
322 95%C.I. (3.95, 15.15), $P = 0.0016$).

323

324 Since these robust models are not accompanied by a model variance it is necessary to also use
325 a mixed effects model system in order to be able to calculate e-Values subsequently. Mixed
326 effects modelling was also conducted after inverse probability weighting. Again a series of
327 increasingly complex models is shown progressing through additive, drug-interactive, full
328 models including drugs, income and ethnicity, and a full model including the two
329 cannabinoids THC and cannabigerol. Importantly in the first three models cannabis is
330 independently highly statistically significant (from β -estimate=79.27 (56.77, 101.78), $P =$
331 1.2×10^{-11}).

332

333 Since the data are gridded in space and time they are well suited for panel linear modelling, a
334 technique which, in addition to inverse probability weighting, allows the added refinements
335 of instrumental variables and temporal lagging. Temporal lagging is pathophysiologically
336 important in such studies as it is likely that any procarcinogenic or environmental exposure
337 takes some time to work before the clinical and epidemiological impact of genotoxicity
338 becomes evident. Again a series of increasingly complex models is presented at increasing
339 lags. Cannabis is again highly significant in many terms, including being independently
340 significant in additive models (from β -estimate=5.31 (1.68, 8.95), $P = 0.0042$).

341

342 Data is also evidently oriented in space and time and is thus eminently suited for formal
343 spatiotemporal analysis. Map-graphs of the data over the 16 years 2002-2017 are shown in
344 Figure 8. Figure 9 shows the geospatial relationships between the contiguous American
345 states and the manner in which links to Hawaii and Alaska have been edited in to define the
346 final spatial neighbourhood network based on “queen” (edge and corner) contiguity. This
347 neighbourhood sparse weights matrix is utilized in all the spatial regressions which follow.

348

349 Table 7 shows the initial results from a series of additive and increasingly complex unlagged
350 interactive spatiotemporal models. The table includes the log of the maximum likelihood

351 ratio (Log.Lik.) at model optimization, and the specifically geospatial model coefficients phi,
352 psi, rho and lambda (see Methods). Since all four of these parameters are generally highly
353 significant this confirms that the full model specification (denoted 'sem2srre' in
354 `splm::spreml`) is appropriate. The Table also lists the standard deviation of each model which
355 is a required input for E-Value calculation. Again cannabis is noted to be independently
356 highly significant in each model.

357

358 Table 8 shows the results of models lagged first just with cannabis and then for all drugs.
359 Interactive terms including cannabis continue to be highly significant. Interactive terms
360 including cannabis are significant from β -estimate=658.72 (396.60, 920.84), $P = 8.40 \times 10^{-7}$
361 for cigarettes: cannabis: alcohol interaction at 2 years of lag.

362

363 Table 9 presents results of models lagged in space for cannabis and in time for the other
364 drugs.

365

366 Table 10 presents the results of temporally lagged interactive space-time models including the two cannabinoids THC and cannabigerol.

367 Cannabigerol is independently significant at 2 lags, and the THC:cannabigerol interaction is significant at zero, two and six lags.

368

369 As mentioned in Methods, well described ethnic disparities exist for many tumours including total cancer. However it is important to consider to

370 what extent such drug use disparities might account for the known epidemiology of TPCIR . Table 11 presents an interactive geospatial

371 regression of the TPCIR against THC exposure of five races as indicated with highly significant results.

372

373 **Table 11.: Spatially- and Temporally- Lagged Spatiotemporal Models**

374

375

376

Parameter			Model				
Parameter	Estimate (C.I.)	P-Value	LogLik	S.D.	Model Parameter	Estimate	P-Value
<i>Cancer Incidence as a Function of Racial Cannabis Exposure</i>							
<i>spreml(Cancer Rate ~ NHWhite THC Exp + NHBlack THC Exp * Hispanic THC Exp * Asian THC Exp * AIAN THC Exp)</i>							
Afric-Am. THC Exp: Hispan.Am THC Exp	1.74 (1.18, 2.29)	1.1E-09	-1532.27	1.9803	phi	0.3887	0.0001
Afric-Am. THC Exp: Hispan.Am THC Exp: Asian-Am. THC Exp: AIAN-Am. THC Exp	0.15 (0.09, 0.21)	1.9E-06			psi	0.1542	0.0005
Asian-Am. THC Exp: AIAN-Am. THC Exp	0.89 (0.37, 1.41)	0.0008			rho	-0.4676	0.0002
Afric-Am. THC Exp: Hispan.Am THC Exp: Asian-Am. THC Exp	-1.11 (-1.55, -0.67)	8.8E-07			lambda	0.4215	8.1E-06
Afric-Am. THC Exp: Hispan.Am THC Exp: AIAN-Am. THC Exp	-0.2 (-0.28, -0.13)	4.8E-08					
Caucasian-American THC Exposure	-1.27 (-1.65, -0.89)	5.0E-11					

377

378

379

380 E-Values are an important way of quantitating the magnitude of co-association required of
381 any unmeasured confounder with both the exposure and outcome variables to explain away
382 the observed effects. Table 12 presents selected E-Value calculations from linear, mixed
383 effects and geospatial models presented in preceding Tables. The key variable to observe is
384 the final number at the right hand side representing the minimum E-Value, and should be
385 read in the light of the observation by one of its originators that E-Values in the literature
386 over approximately 1.25 are considered noteworthy [56]. In general terms the E-Values fall
387 in the sequence geospatial models > mixed effects models > linear models, related partly to
388 the much smaller model variance of more complex models.

389

390 Table 12 lists 56 E-Values related to cannabis or cannabinoids of which 24 are larger than
391 1,000. Of the 33 E-Values originating from geospatial models, 20 are larger than 1,000. The
392 table lists six minimum e-Values of infinity, three deriving from mixed effects models and
393 three from geospatial models.

394

395 Given the above compelling data demonstrating a link between rising rates of cannabis
396 exposure and rising TPCIR an obvious extension of this study was whether the increasing
397 use, availability and concentration of cannabis associated with more liberal legal paradigms
398 [57] was associated with elevated TPCIR. One important caveat on such an investigation is
399 that since the data only run to 2017 and many populous states had not yet been affected by
400 the cannabis legalization movement, it may be considered that the data is premature for a full
401 determination of this potential effect. Figure 10A shows the rate of TPCIR under various
402 legal paradigms. Whilst the few states involved with full cannabis legalization at that time
403 were associated with broad confidence interval bands there is a clear impression in this
404 Figure that the rate under decriminalization appeared to be at a higher levels than others.
405 Figure 10B dichotomizes the data into liberal paradigms vs. traditional policies of cannabis
406 being considered illegal. Separation of the two regression lines towards the right hand side of
407 the graph gives a clear impression for a significant interaction between time and
408 dichotomized legal status.

409

410 These differences are formally assessed in Table 13 by linear regression. Decriminalized and
411 legal status are both confirmed to be significant on their own (upper table segment). In
412 interaction with time decriminalized status is significant (middle table segment).
413 Dichotomized legal status is also found to be significant in interaction with time (lower table

414 segment, β -estimate= 1.87×10^{-4} , (2.9×10^{-5} , 2.45×10^{-4}), $P=0.0208$). Table 12 lists the minimum
415 E-Values associated with these changes as 1.60 and 1.98 for cannabis decriminalization and
416 full cannabis legalization respectively (at the bottom of the Linear Regression part of Table
417 12).
418

419 Discussion

420

421

422 Main Results

423

424 The main results of this study confirmed that total Pediatric cancer rates have risen
425 significantly nationally across USA and this trend holds for the commonest pediatric
426 malignancies the leukaemias, Non-Hodgkins lymphoma, localized and distant sarcoma and
427 testicular cancer. It was important to note across this period that the use of tobacco, alcohol
428 use disorders, cocaine and analgesic abuse declined as measured in major national surveys
429 whilst cannabis use alone was rising. The level of cannabinoids identified in Federal seizure
430 data also rose for most cannabinoid analytes. TPCIR rose strongly and significantly as a
431 function of cannabinoid exposure, but only weakly and non-significantly in bivariate analysis
432 in relation to cannabis itself. TPCIR was significantly higher in the two highest cannabis use
433 quintiles both overall and across time. Inverse probability weighting was used to equilibrate
434 cannabis exposure across the cohort. Indices of ethnic cannabinoid exposure and seizure
435 cannabinoid concentrations were variously used as instrumental variables to adjust panel
436 models.

437

438 Cannabis use was independently associated with TPCIR in additive robust marginal
439 structural, mixed effects, panel and geospatiotemporal models. Cannabis use was
440 independently associated with TPCIR in interactive mixed effects and geospatial models.
441 Cannabis use was linked with TPCIR in various interactions in linear models, robust
442 marginal, mixed effects, panel and geospatial models. Cannabis was independently linked
443 with TPCIR in geospatial models lagged to zero, 1 and 6 years and featured in interactions
444 lagged to 1,2,4 and 6 years. When the cannabinoids THC and cannabigerol were studied they
445 were also linked with TPCIR at high levels of statistical significance at zero, 2, 4 and six
446 years of lag.

447

448 On sensitivity analysis 49 of 56 minimum e-Values were above 1.25 which is a quoted
449 threshold for likely causal relationships. Similarly 31 of 33 geospatial e-Values were above
450 this threshold. The highest finite minimum e-Value was 4.14×10^{89} . Six minimum e-Values
451 were infinity.

452

453 The recent trend to cannabis liberalization was associated with elevated TPCIR both as a
454 group and as an acceleration of the time-dependent trend in cannabis-liberal states.

455

456 Our interpretation of these highly consistent and concordant findings obtained by several
457 methodologies with instrumental variables, controlling for ethnic cannabinoid exposure,
458 utilizing robust regression techniques, inverse probability weighting with high levels of
459 association across both space and time together with very high e-Values is that the
460 relationship of cannabinoid exposure to total pediatric cancer incidence fulfills the criteria of
461 causality and explains the increasing rates of pediatric cancer under cannabis-liberal
462 legislative paradigms, and that this statement is especially true for THC and cannabigerol the
463 two cannabinoids which show the most consistent rises over time.

464

465 Hence our study is closely concordant with other published series on the link between
466 pediatric cancer and cannabis use [7-11].

467

468

469 Statistical Comments

470 It is worth considering briefly the incisive logical power of space-time regression. To say
471 that two variables are statistically associated carries a certain weight. To say that two
472 variables are closely associated when their distribution is considered across both space and
473 time simultaneously is strongly suggestive of a presumptively causal relationship.

474

475 Nineteen spatiotemporal models were presented. In seventeen the spatial error coefficient
476 rho was significant. In eighteen the spatial error autocorrelation coefficient lambda was
477 significant. And spatial errors adjusted in the manner of Kapoor, Kelejian and Prucha
478 consistently had higher precision than those adjusted by the algorithm of Baltagi. Together
479 this is indisputable evidence of effects operating in a spatially distributed manner, and
480 represents in the data analytical environment a reflection of the orchestrated campaign across
481 USA to legalize cannabis which operated in a coordinated manner from the west coast
482 eastwards.

483

484 Some comments in relation to casual inference and causal assignment are pertinent. Inverse
485 probability weighting is a method which is well established to correct for inconsistent
486 exposures amongst groups. It enjoys a strong theoretical and epidemiological evidence

487 base [58]. Its effect is to even out exposures between groups and creates pseudo-randomized
488 populations from which causal implications can appropriately be drawn. Similarly E-Values
489 were recently introduced in a formal way to quantitate extraneous confounding from
490 unmeasured covariates and provides a quantitative magnitude to the level of association
491 required of unknown factors with both the exposure and the outcome to remove the impact of
492 any described association [59].

493

494

495

496 Mechanisms

497

498

499 Of pivotal importance in linking associational findings with causal pathways is the issue of
500 biological plausibility and the cellular and molecular pathways which might connect the
501 exposure of interest with the outcome of concern. The subject of the pro-oncogenic activities
502 and potential of cannabis, cannabis smoke and cannabinoids is complex major papers have
503 addressed this issue [14, 26, 28, 32, 34-36, 42, 60-66]. In this paper we will provide a brief
504 and concise overview of what presently seem to be some of the most important pathways
505 which are likely to be implicated. They will be described under nine headings of:
506 gametotoxicity, genotoxicity, epigenotoxicity, mitochondriopathy, immunomodulation, pro-
507 aging, endovascular ischaemia – hypoxia, sympathetically mediated effects on stem cell
508 niches and non-linearity of the dose-response genotoxic effect curve. These domains are not
509 independent but are themselves interdependent and intricately intertwined. Whilst most of
510 the following observations have been experimentally defined the logical sequence has been
511 filled out where this seems reasonable and concordant with the evidence base.

512

513 Cannabinoids have been detected in seminal fluid and have been linked with DNA nicking
514 and fragmentation, abnormal sperm nuclear size, gross abnormalities of sperm morphology
515 including sperm fragmentation, disordered DNA packing and re-packing, disorders of
516 protamine synthesis, histone-protamine substitution and major disruption of sperm DNA
517 methylation [15-17, 31, 37, 61, 67, 68]. Cannabinoids have been found in Graafian follicle
518 and oviduct fluid and have been linked with oocyte nuclear blebbing, nuclear bridging,
519 chromosomal fragmentation and large scale oocyte loss after the second meiotic cell division
520 [14, 15, 17]. Cannabis smoke is known to contain all of the carcinogens of tobacco smoke

521 including many tars and carcinogens including aromatic amines, polycyclic hydrocarbons,
522 and tars [69]. Cannabinoid exposure has been linked with nuclear bleb and chromosomal
523 bridge formation, chromosomal mis-segregation at the anaphase separation, micronucleus
524 formation [70], transposon activation and chain and ring chromosome formation [14, 32, 34].
525 Cannabidiol, Cannabinol and THC have been implicated in in chromosomal translocation
526 formation to the same level seen with cytotoxic drugs [13]. Cannabidiol and cannabidivarin
527 have been shown to cause double stranded DNA breaks, micronucleus formation and nuclear
528 buds and bridges in human cells which is worse under oxidative stress [66]. Cannabinoid-
529 induced micronucleus formation is very important as it has been identified as a major engine
530 of catastrophic damage to the genetic material and one-step chromothripsis, chromoanagenesis
531 and oncogenic transformation [60, 71, 72]. Cannabinoid exposure has been linked with large
532 scale perturbation of DNA methylation, gross defects in histone synthesis – which necessarily
533 leave DNA more open and available for transcription which is a pro-oncogenic state – altered
534 histone signalling, and an inhibition of ATP supply to genetic and epigenetic processes –
535 most of which are energy dependent – and an inhibition of epigenetic substrate supply [31,
536 33, 35, 37, 61, 73]. Together these changes may be expected to advance the “epigenetic
537 clock” which is believed to be one of the key determinants of cellular aging [74, 75]. The
538 profound implications of major epigenetic reprogramming were highlighted by a recent paper
539 noting that despite the short half life of immune cells in the circulation – just a few days - the
540 cellular basis for long lasting immunity is actually epigenetic changes in long lived myeloid
541 precursor cells which record metabolic and immune activation responses in the coordinated
542 patterns of their enhancers, promoters, long non-coding RNA’s, DNA methylation and
543 histone codes which determine chromatin conformation and the assembly of topologically
544 transcriptionally active domains which functionally facilitate secondary responses to
545 infection and vaccines [76, 77].

546

547 The outer mitochondrial membrane not only possess CB1R’s, but indeed the whole of the
548 cannabinoid signalling transduction machinery found in the plasmalemma also resides in the
549 inner and outer mitochondrial membrane and within the intermembrane space so that
550 cannabinoids are an important direct modulator of metabolic state [78-82]. Several adverse
551 mitochondrial processes are well described including a reduction in the transmembrane
552 potential across the inner mitochondrial membrane, a reduced synthesis of key oxidative
553 phosphorylation substrates including the F1-ATPase, increased electron shunting via

554 uncoupling protein 2 activation, gross mitochondrial damage and swelling and impairment of
555 mitonuclear cross-talk and mitonuclear genomic coordination [17, 83-87].

556

557 There is a rich literature describing both the pro- and anti- inflammatory actions of
558 cannabinoids. In this context the proinflammatory CB1R-mediated activities seem to be
559 especially important [88] as chronic inflammation is a well established cause of cancers in
560 many tissue beds and occurs by many mechanisms. One pathway of particular interest is that
561 cytoplasmic inflammation stimulates the transposons or “jumping genes” of the genome, to
562 start “jumping” mobile segments and creating genomic havoc. Micronucleus disruption
563 releases double stranded DNA into the cytoplasm where it potently stimulates the
564 cytoplasmic GMP-AMP – Stimulator of Interferon Gamma (cGAS-STING) pathway which
565 further intracytoplasmically stimulates inflammation via interferon- γ and innate immune
566 signalling and destabilizes the genome [89-91]. The immunosuppressive activities of
567 cannabinoids may depress the immune response to the developing field change and nascent
568 tumours. This cycle could potentially explain the many case reports of cancers occurring in
569 adults at a younger age than usual and with increased aggressiveness in heavily cannabis
570 exposed patients [92-95].

571

572 Cannabis exposure has been found to accelerate organismal cardiovascular aging clinically
573 [96]. Cannabinoids are known to inhibit stem cell division [34, 97]. This combination of
574 impaired stem cell activity, reduction of mitochondrial energy generation and a pro-
575 inflammatory milieu are all hallmarks of cellular ageing and the senescence-associated
576 secretory phenotype [98-100] of growth factors and cytokines which is presumably
577 stimulated and a key hallmark of aging. Aging of course is the leading risk factor for most
578 adult tumours. In the light of the foregoing cellular changes it would seem that the quality of
579 cannabinoid-exposed gametes may be broadly seen as defective and they may thus be said in
580 general terms to likely be “aged” in metabolic, epigenetic and genetic terms. Cannabinoids
581 are known to have important effects on the microvasculature and can induce tissue ischaemia
582 [101-104] which is an important determinant of the hypoxic microenvironment which
583 stimulates genomic instability and oncogenesis and promotes nascent and mature tumour
584 growth. Cannabis addiction is known to feature periods of cannabinoid withdrawal marked
585 by agitation and manifest sympathetic hyperstimulation [105]. Sympathetic stimulation has
586 been shown to have direct adverse activities on the stem cell niche of the hair follicle [106]
587 and likely acts similarly in other stem cell niches.

588

589 Arguably the most concerning feature of this literature is the apparent threshold effect beyond
590 which genotoxic and mitochondriopathic changes emerge relatively abruptly. This implies
591 that the exponential dose-effect curve seen in many genotoxic assays for cannabinoids [35,
592 62, 107, 108] can appear to be functionally an abrupt discontinuity in the dose-response curve
593 at the epidemiological level. At the community level this implies that a doubling of daily
594 cannabis use, as has been documented in USA in recent years [109], might reasonably be
595 linked with a disproportionate response in genotoxic downstream sequelae such as
596 congenital anomalies including transgenerationally transmissible carcinogenesis.

597

598 From this brief overview it is apparent that a plethora of cellular oncogenic mechanisms exist
599 linking exposure to cannabis smoke, cannabis and cannabinoids to the processes of
600 carcinogenesis.

601

602

603 In 1965 Hill described nine criteria as being required of any association in order to assign
604 causality to the relationship. Strength of association, consistency amongst studies,
605 specificity, temporal sequence, coherence with known data, biological plausibility a
606 biological response or dose-response curve, analogy with similar situations elsewhere and
607 experimental confirmation were key features [110]. It will be noted that the above analysis,
608 including the published literature and the cited experimentally demonstrated mechanistic
609 links, fulfill all of these criteria for the relationship between cannabis exposure and TPCIR .

610

611

612 Generalizability

613

614 Our data are population level data derived from publicly available datasets from one of the
615 world's most technologically advanced nations. The underlying population is also
616 substantial. Given that our findings are robust to various different methods, fulfill criteria for
617 causality and are consistent with the majority of the published work in the area we believe
618 that our findings are robust and widely generalizable. However as it is clear that cannabis use
619 is in a state of flux worldwide at the present with rises in the prevalence of use, intensity of
620 use, and concentration of product we feel that it is important that on-going studies be
621 conducted in this area to monitor the situation at higher levels of geospatial resolution.

622

623

624 Future Directions

625

626 Further extensions of this work might include detailed dissection of the molecular and
627 cellular level of the pathways mentioned particularly relating to mitochondrial cannabinoid
628 signalling, mitochondrial electron leaks and shunts, free oxyradical flux, perturbation of
629 mitonuclear cross-talk, cannabinoid induced disruption of metabolic supply of epigenetic
630 substrates, cannabinoid-related disruption of histone synthesis and signalling and the histone
631 code generally, cannabinoid epigenotoxicity generally and heritable and transgenerational
632 epigenotoxicity specifically, proinflammatory cannabinoid actions, microvascular-disrupting
633 and hypoxia-inducing actions, chromosomal mis-segregation and anaphase disruption and the
634 interaction of cannabinoids with the cGAS-STING cytoplasmic signalling pathway.

635 Research into cannabinoid interactions with the germ cells, oocytes and sperm, is clearly of
636 primary and foundational importance to these concerns and should be up-prioritized on
637 research agendas. Analytically higher resolution space-time modelling based on more
638 detailed datasets from CDC and SAMHSA is an obvious task for the near future. The
639 incorporation of instrumental variables and inverse probability weights into the space-time
640 and spatiotemporally lagged models of plm, splm and similar software would allow all the
641 questions of interest to be addressed in a single modelling framework without the need for
642 multiple model types as was necessitated in the present report and would likely only require
643 minimal resources to enable the required programming code to be written for this very
644 impressive, sophisticated and highly flexible software to be further optimized.

645

646

647 Strengths and Limitations.

648

649 Our study has several strengths including using data from a very populous nation, the use of
650 publicly available datasets, the use of different statistical techniques, the application of
651 inverse probability weighting and e-Values, two mechanisms well established in the causal
652 epidemiological literature, the use of geospatiotemporal regression techniques with complex
653 random error structures, the use of models lagged both spatially and temporally, the use of a
654 variety of covariates, consideration of substance-exposure indices which is often absent from
655 many studies, the use of various instrumental variables, the availability of a relatively lengthy

656 panel data series for 15 years, and correction for ethnic cannabis exposure as a major
657 underlying confounding factor. The absence of geospatial techniques from much of cancer
658 epidemiology appears to be a major knowledge gap which the present study begins to redress.
659 It may also be argued that for research enterprises to consume significant public resources but
660 never be able to provide actual causal advice to their host community at once stretches public
661 credulity and tests their patience, particularly when well established methodologies are
662 available which can be used to fill this major knowledge gap. The deliberate application of
663 the techniques of formal causal inference in this study thus comprises a major strength. The
664 study's major limitation relates to the unavailability of individual patient-level data which is a
665 common limitation amongst epidemiological studies. Due to the complexity of the present
666 analysis we have not considered further subgroup analyses, either of individual tumours, or
667 by fascinating sex or ethnic incidence differences. All of this remains to be done at higher
668 geospatial resolution by subsequent investigators.

669

670

671 Conclusion

672

673 In summary our study confirms previous reports in the literature linking cannabis exposure
674 with pediatric and testicular cancer [7-11, 18-22] and answers both our opening hypotheses
675 affirmatively. We extend and amplify earlier reports in many ways including with the use of
676 national cancer census data and widely cited nationally representative drug use surveys, the
677 application of geospatial techniques and the formal techniques of causal inference to the data
678 series and various technical refinements including the use of several sets of instrumental
679 variables and various forms of inverse probability-weighted and spatially weighted regression
680 matrices and robust, panel and linear multivariable techniques. After including
681 socioeconomic, ethnic and drug use variables we find robust associations across space and
682 time for cannabis use and TPCIR and that cannabis, and particularly the cannabinoids THC
683 and cannabigerol, are independently and interactively associated with TPCIR both in *de novo*
684 space-time grids and in spatially and temporally lagged models. Moreover very high e-
685 Values clearly indicate that the relationship cannot be explained away by unmeasured,
686 unknown or hypothetical confounding variables. This analysis is consistent with five
687 previously reported series comprising the majority of the published literature in the field [7-
688 11], dozens of potential experimentally described mechanistic pathways and fulfill the
689 paradigmatic Hill criteria of causality [110]. Findings are also consistent with reports of

690 elevated rates of congenital anomalies following prenatal cannabis exposure [25-28, 42, 43]
691 and thus are broadly concordant conceptually with wide ranging and far reaching heritable
692 cannabinoid-related genotoxicity. Our analysis also begins to provide insights into the
693 previously mysterious major differences in cancer incidence between various ethnicities by
694 indicating that varying ethnic exposures to cannabinoids are of particular concern. It is
695 important that this thread be further explored in the future. Such formal demonstration of
696 strong evidence of a presumptively genotoxic cannabis-cancer causal link is highly relevant
697 for the ongoing and currently controversial story of the relationship of cannabis use with
698 malignant tumourigenesis in adults. Strong evidence of a robust causal relationship of
699 cannabis exposure to pediatric and thus transgenerational inheritable genotoxicity carries far
700 reaching implications for the ongoing public debate relating to the most appropriate forms of
701 regulation of cannabis and cannabinoids. Moreover the present analysis powerfully informs
702 the broader discussion regarding cannabis-related genotoxicity as it relates to adult
703 tumourigenesis and many congenital anomalies encountered at birth [25-28, 42, 60, 61].

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708 Declarations

709

710 Ethics Approval and Consent to Participate

711 The Human Research Ethics Committee of the University of Western Australia provided

712 ethical approval for the study to be undertaken 7th June 2019 (No. RA/4/20/4724). Ethics

713 approval was not required to access the data in the first instance. However Ethical approval

714 provided permission to access, analyze and publish all the data obtained.

715

716 Consent for Publication

717 Not applicable.

718

719 Availability of Data and Materials

720 All data generated or analysed during this study are included in this published article and its

721 supplementary information files. No permissions are required to access the data which was

722 used and collated in this study, e.g. NSDUH study. Data including shapefiles and R

723 programming script is made publicly available on the Mendeley Data Archive at this URL:

724 URL: <http://dx.doi.org/10.17632/cnww9hdspd.1> .

725

726

727 Competing Interests

728 The authors declare that they have no competing interests.

729

730

731 Funding

732 No funding was provided for this study. No funding organization played any role in the

733 design and conduct of the study; collection, management, analysis, and interpretation of the

734 data; preparation, review, or approval of the manuscript; and decision to submit the
735 manuscript for publication.

736

737

738 Authors' Contributions

739 ASR assembled the data, designed and conducted the analyses, and wrote the first manuscript
740 draft. GKH provided technical and logistic support, co-wrote the paper, assisted with gaining
741 ethical approval, provided advice on manuscript preparation and general guidance to study
742 conduct.

743

744 Acknowledgements

745 Not applicable.

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Table 1.: SEER-Nominated Time Trends of Various Pediatric and Adult Cancers

Cancer	Observed Trend	Delayed Trend
All Pediatric Cancers (<20 Years)	Rising	Rising
Pediatric ALL - Acute Lymphatic Leukaemia	Rising	
Pediatric AML - Acute Myeloid Leukaemia	Rising	
Pediatric Brain Cancer	Stable	Rising
Pediatric NHL - Non-Hodgkins Lymphoma	Rising	Rising
Sarcoma - All Age	Stable	
Sarcoma <20 Year - Localized	Rising	
Sarcoma <20 Year - Distant	Rising	
Sarcoma All Age - Localized	Rising	
Sarcoma All Age - Distant	Rising	
Sarcoma All Age	Rising	
Pediatric Testes < 20 Years	Stable	Stable
Testes < 50 Years	Rising	Rising
Testes All Age	Rising	Rising

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Table 2.: Linear Models: TPCIR Against Time, Cannabis, Cannabinoids and Ethnicity

Parameter Estimates			Model Parameters			
Parameter	Estimate (C.I.)	Pr(>t)	R-Squared	F	dF	P
<i>lm(Cancer_Rate ~ Time)</i>						
Year	0.14 (0.1, 0.17)	3.8E-14	0.0725	59.6	1,748	3.80E-14
<i>lm(Cancer_Rate ~ Cannabis)</i>						
mrjmon	1.00 (-1.22, 3.22)	0.3800	-0.0003	0.78	1,748	0.3770
<i>lm(Cancer_Rate ~ Δ9THC)</i>						
Δ9THC	0.33 (0.15, 0.5)	0.0002	0.0169	13.8	1,748	0.0002
<i>lm(Cancer_Rate ~ Exposure * Drug)</i>						
Drug_Rate: Cannabis	4.63 (2.11, 7.15)	0.0003	0.0207	9.82	9,3740	5.39E-15
Drug_Rate: Alcohol	-3.22 (-6.21, -0.22)	0.0356				
Drug_Rate: Analgesics	-6.63 (-10.51, -2.75)	0.0008				
Cocaine	-1.06 (-1.63, -0.49)	0.0003				
Cannabis	-1.32 (-1.89, -0.74)	0.0000				
Drug_Rate	-3.63 (-4.86, -2.4)	0.0000				
<i>lm(Cancer_Rate ~ Exposure * Cannabinoid)</i>						
Cannabinol	6.54 (5.07, 8.01)	< 2E-16	0.0402	18.9	7,2992	<2E-16
Cannabigerol	7.65 (5.91, 9.38)	< 2E-16				
Drug_Rate	2.14 (1.55, 2.73)	0.0000				
Cannabichromene	3.86 (0.29, 7.42)	0.0340				

Drug_Rate: Cannabichromene	-3.02 (-5.4, -0.63)	0.0130				
<i>lm(Cancer_Rate ~ Ethnic_THC_Exposure * Ethnicity)</i>						
Ethnic_THC_Exposure	0.14 (0.07, 0.21)	0.0001	0.0021	2.57	6,4493	0.0174
Asian-Am_THC_Exposure	0.28 (0.02, 0.55)	0.0360				

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Table 4.: Robust Generalized Linear Regression Models1088
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Parameter	Estimate (C.I.)	P-Value
<i>Additive Model</i>		
<i>svyglm(Cancer_Rate ~ Cigarettes + Cannabis + Analgesics + Alcohol + Cocaine)</i>		
Cannabis	9.55 (3.95, 15.15)	0.0016
Alcohol	-19.69 (-27.68, -11.7)	1.5E-05
<i>Interactive Model</i>		
<i>svyglm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine)</i>		
Cigarettes: Cannabis: Analgesics	268.42 (91.87, 444.96)	0.0046
Cigarettes: Analgesics	-59.54 (-92.24, -26.84)	0.0009
<i>Full Interactive Model</i>		
<i>svyglm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine + 6_Races + Income)</i>		
White	8.1 (6.04, 10.17)	4.2E-09
Hispanic	0.74 (0.37, 1.11)	0.0004
Asian	0.77 (0.38, 1.16)	0.0004
Cigarettes: Alcohol: Analgesics	331.59 (121.58, 541.61)	0.0038
Cigarettes: Cannabis: Analgesics	2537.45 (833.94, 4240.95)	0.0060
Cigarettes	52.94 (15.62, 90.27)	0.0086
Alcohol: Analgesics	871 (196.96, 1545.04)	0.0158
Cannabis: Alcohol	543.49 (110.37, 976.6)	0.0189
Cigarettes: Cannabis	-268.79 (-471.82, -65.76)	0.0136
Alcohol	-119.12 (-207.37, -30.87)	0.0120
Cannabis: Alcohol: Analgesics	-4989.69 (-8616.76, -1362.61)	0.0106

AIAN	-6.66 (-11.36, -1.95)	0.0087
Cigarettes: Analgesics	-500.17 (-808.72, -191.63)	0.0030
Full Interactive Model with Cannabinoids		
<i>svyglm(Cancer_Rate ~ Cigarettes * Δ9THC * Cannabigerol * Alcohol + Analgesics + Cocaine + 6_Races + Income)</i>		
White	7.87 (5.73, 10.02)	1.9E-08
Cocaine	25.98 (12.75, 39.21)	0.0005
Asian	0.68 (0.31, 1.06)	0.0010
Hispanic	0.59 (0.23, 0.94)	0.0026
Cigarettes: Δ9THC: Analgesics	34.32 (13.53, 55.11)	0.0026
Cigarettes: Cannabigerol	270.35 (104.12, 436.59)	0.0030
Cigarettes: Δ9THC	2.93 (0.58, 5.28)	0.0195
Δ9THC: Cannabigerol	29.24 (4.95, 53.53)	0.0239
AIAN	-5.96 (-11.16, -0.75)	0.0311
Cigarettes: Δ9THC: Alcohol	-13.34 (-24.92, -1.77)	0.0300
Cigarettes: Δ9THC: Cannabigerol	-103.55 (-181.75, -25.34)	0.0136
Cannabigerol	-115.2 (-189.34, -41.06)	0.0043
Cigarettes: Analgesics	-87.59 (-127.51, -47.66)	1.2E-04

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Table 5.: Mixed Effects Regression Models

Parameters			Model Parameters			
Parameter	Estimate (C.I.)	P-Value	SD	AIC	BIC	logLik
<i>Additive Model</i>						
<i>lme(Cancer_Rate ~ Cigarettes + Cannabis + Analgesics + Alcohol + Cocaine)</i>						
Cannabis	5.34 (0.07, 10.6)	0.0472	3.43138	3884.77	3912.46	-1936.39
Analgesics	-11.02 (-18.65, -3.39)	0.0048				
<i>Interactive Model</i>						
<i>lme(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine)</i>						
Cannabis	72.88 (49.6, 96.15)	1.4E-09	3.31033	3781.12	3836.4	-1878.56
Cigarettes	43.36 (27.68, 59.04)	8.2E-08				
Alcohol: Analgesics	1523.99 (970.61, 2077.38)	9.3E-08				
Cigarettes: Cannabis: Analgesics	2788.19 (1676.17, 3900.2)	1.1E-06				
Cannabis: Alcohol: Analgesics	-4554.93 (-6709.17, -2400.69)	3.8E-05				
Cigarettes: Analgesics	-539.08 (-790.18, -287.99)	2.9E-05				
Analgesics	-87.43 (-121.63, -53.23)	6.9E-07				
Alcohol	-82.06 (-113.58, -50.54)	4.3E-07				
Cigarettes: Cannabis	-284.5 (-376.55, -192.45)	2.3E-09				
<i>Full Interactive Model</i>						
<i>lme(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine + 6 Races + Income)</i>						
White	11.8 (8.45, 15.14)	1.1E-11	3.18221	3715.57	3784.61	-1842.79
Cannabis	79.27 (56.77, 101.78)	1.2E-11				
Asian	2.54 (1.8, 3.27)	2.6E-11				
Cigarettes: Alcohol: Analgesics	1636.35 (1108.24, 2164.46)	2.1E-09				

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Cigarettes	45.74 (30.44, 61.04)	7.2E-09				
Cigarettes: Cannabis: Analgesics	2525.7 (1488.65, 3562.75)	2.2E-06				
Alcohol: Analgesics	959.4 (425.8, 1493)	4.5E-04				
Cannabis: Alcohol: Analgesics	-4264.85 (-6314.08, -2215.61)	5.1E-05				
Alcohol	-93.44 (-124.44, -62.43)	5.5E-09				
Cigarettes: Analgesics	-766.56 (-1011.12, -521.99)	1.4E-09				
Cigarettes: Cannabis	-290.63 (-373.83, -207.42)	1.7E-11				
Income	-9.44 (-12.02, -6.87)	1.7E-12				
Full Interactive Model with Cannabinoids						
$\ln(\text{Cancer_Rate} \sim \text{Cigarettes} * \Delta\text{9THC} * \text{Cannabigerol} * \text{Alcohol} + \text{Analgesics} + \text{Cocaine} + 6 \text{ Races} + \text{Income})$						
White	15.39 (11.82, 18.96)	1.8E-16	3.16296	3743.28	3798.56	-1859.64
Asian	2.46 (1.76, 3.16)	1.2E-11				
Cigarettes: Cannabigerol: Alcohol	4741.19 (3077.86, 6404.51)	3.3E-08				
Cigarettes: Δ9THC	26.57 (15.54, 37.6)	2.8E-06				
Δ9THC : Alcohol	14.95 (7.74, 22.16)	5.4E-05				
Hispanic	0.7 (0.14, 1.26)	1.4E-02				
Cigarettes: Cannabigerol	-663.69 (-971.24, -356.13)	2.7E-05				
Income	-7.76 (-10.11, -5.41)	1.9E-10				
Cigarettes: Δ9THC : Alcohol	-240.65 (-304.57, -176.72)	4.6E-13				

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1113**Table 6.: Panel Regression Models**

Model Specification		Parameters			Model Parameters				
Instrumental Variables	Lagged Parameter	Parameter	Estimate (C.I.)	P-Value	Adj. R-Squared	Chi.S qu.	F	dF	P
		<i>Additive model</i>							
		<i>plm(Cancer_Rate ~ Cigarettes + Cannabis + Analgesics + Alcohol + Cocaine)</i>							
		Cannabis	5.31 (1.68, 8.95)	0.0042	0.0790	80.6858		3	<2.2E-16
		Analgesics	-9.3 (-14.93, -3.67)	0.0012					
		Cigarettes	-4.53 (-7.15, -1.92)	0.0007					
		<i>Interactive model</i>							
		<i>plm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine)</i>							
		Cigarettes: Cannabis	24.47 (11.37, 37.57)	0.0003	0.0663	83.1987		4	<2.2E-16
		Cocaine	-11.93 (-22.84, -1.02)	0.0321					
		Analgesics	-8.25 (-14.2, -2.3)	0.0066					
		Cigarettes	-5.83 (-8.92, -2.73)	0.0002					
		<i>Interactive Full model</i>							
		<i>plm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine + 6_Races + Income)</i>							
		White	8.57 (7.18, 9.96)	<2.2E-16	0.1927	23.5694		13,736	<2.2E-16
		Asian	0.92 (0.67, 1.17)	2.0E-12					
		Hispanic	0.71 (0.47, 0.94)	7.1E-09					
		Cigarettes	69.38 (39.16, 99.59)	7.9E-06					
		Cigarettes: Alcohol: Analgesics	1169.92 (598.17, 1741.68)	6.7E-05					
		Cannabis: Alcohol	719.64 (352.45, 1086.83)	1.3E-04					

		Cigarettes: Cannabis :Analgesics	2926.99 (1407.72, 4446.25)	0.0002					
		Analgesics	58.43 (22.7, 94.17)	0.0014					
		Alcohol: Analgesics	709.96 (139.61, 1280.31)	0.0149					
		Cannabis: Alcohol: Analgesics	-5916.9 (-9125.97, -2707.83)	0.0003					
		Cigarettes: Cannabis	-345.82 (-521.51, -170.13)	1.2E-04					
		Alcohol	-153.51 (-219.89, -87.13)	6.8E-06					
		Cigarettes: Analgesics	-716.53 (-1007.91, -425.15)	1.8E-06					
		Interactive Full model - 2 Lags							
		plm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine + 6_Races + Income)							
	Cigarettes, 2	White	9.28 (7.88, 10.67)	<2.2E-16	0.2014		33.9397	7,642	2.0E-01
	Cannabis, 2	Asian	0.96 (0.71, 1.22)	2.1E-13					
	Analgesics, 2	Hispanic	0.67 (0.41, 0.94)	9.4E-07					
	Alcohol, 2	Cigarettes	18.29 (9.29, 27.29)	7.6E-05					
	Cocaine, 2	Cigarettes: Alcohol: Analgesics	513.89 (96.6, 931.17)	0.0161					
		Cigarettes: Analgesics	-92.64 (-154.24, -31.03)	0.0033					
		Cigarettes: Alcohol	-95.72 (-146.79, -44.65)	0.0003					
		Interactive Full model - 4 Lags							
		plm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine + 6_Races + Income)							
	Cigarettes, 4	White	8.71 (6.97, 10.46)	<2.2E-16	0.1990		17.6055	12,537	<2.2E-16
	Cannabis, 4	Hispanic	0.85 (0.54, 1.16)	7.3E-08					
	Analgesics, 4	Asian	0.72 (0.4, 1.05)	1.7E-05					
	Alcohol, 4	Cigarettes: Cannabis	233.66 (115.14, 352.18)	1.3E-04					
	Cocaine, 4	Cigarettes: Alcohol: Analgesics	1975.86 (855.19, 3096.52)	0.0006					
		Cannabis: Alcohol: Analgesics	1972.91 (573.99, 3371.83)	0.0059					
		Alcohol	105.37 (17.01, 193.73)	0.0198					
		AIAN	-8.33 (-14.85, -1.8)	0.0127					
		Alcohol: Analgesics	-647.54 (-1104.4, -190.68)	0.0057					
		Cannabis: Alcohol	-376.97 (-619.06, -134.87)	0.0024					
		Cigarettes: Alcohol	-286.48 (-462.48, -110.48)	0.0015					
		Cigarettes: Cannabis: Analgesics	-1300.25 (-2003.68, -596.83)	0.0003					

		Interactive Full model - 5 Lags							
		plm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine + 6_Races + Income)							
THC	Cigarettes, 5	White	6.7228 (5.29, 8.15)	<2.2E-16	0.2302	233.988		10	<2.2E-16
Cannabigerol	Cannabis, 5	Hispanic	0.652 (0.41, 0.89)	7.2E-08					
Cannabinol	Analgesics, 5	Asian	0.6755 (0.42, 0.93)	2.6E-07					
Cannabichrome ne	Alcohol, 5	Cigarettes: Cannabis	28.717 (14.28, 43.16)	0.0001					
	Cocaine, 5	Cigarettes: Cannabis: Alcohol: Analgesics	8122.4789 (3192.57, 13052.39)	0.0012					
		Cannabis: Analgesics	463.7354 (130.9, 796.57)	0.0063					
		Cigarettes: Cannabis: Analgesics	-1099.2482 (-1927.82, -270.68)	0.0093					
		Cannabis: Alcohol: Analgesics	-3523.9404 (-5495.28, -1552.6)	0.0005					
		AIAN	-9.4157 (-14.03, -4.81)	0.0001					
		Cigarettes	-5.1117 (-7.44, -2.78)	1.7E-05					
		Interactive Full Model with Racial Cannabis Exposure as Instrumental Variables							
THC Exposure		plm(Cancer_Rate ~ Cigarettes * Cannabis * Analgesics * Alcohol + Cocaine + 6_Races + Income)							
In:		White	0.7294 (-11.72, 25.62)	<2.2E-16	0.2300	232.721		9	<2.2E-16
Caucas-Am.		Asian	0.1306 (-9.77, 11.17)	9.2E-08					
African-Am.		Hispanic	0.1202 (-9.65, 10.91)	1.6E-07					
Hispan-Am.		Cigarettes: Cannabis: Alcohol: Analgesics	731.4135 (3112.74, 3129.47)	2.0E-05					
Asian-Am.		Cannabis: Analgesics	35.5327 (136.48, 152.42)	4.8E-05					
AIAN-Am.		Cocaine	4.9629 (7.98, 18.41)	0.0078					
NHPI-Am.		AIAN	2.3266 (-1.45, -16.93)	7.9E-05					
		Cannabis: Alcohol: Analgesics	378.1363 (-1752.55, -1770.82)	3.2E-06					
		Cigarettes: Analgesics	9.4453 (-37.01, -56.39)	7.7E-07					

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Table 7.: Introductory Spatiotemporal Models1116
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Parameter			Model				
Parameter	Estimate (C.I.)	P-Value	LogLik	S.D.	Model Parameter	Estimate	P-Value
<i>Additive Model</i>							
<i>spreml(Cancer_Rate ~ Cigarettes + Cannabis + Alcohol + Analgesics + Cocaine)</i>							
Cannabis	5.16 (2.26, 8.06)	0.0005	-1541.00	1.9451	phi	0.3170	0.0002
Analgesics	-4.6 (-9.18, -0.02)	0.0490			psi	0.1480	0.0007
Cigarettes	-2.72 (-4.85, -0.59)	0.0124			rho	-0.4959	2.2E-05
					lambda	0.4598	8.2E-08
<i>3-Way Interactive model</i>							
<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol + Analgesics + Cocaine)</i>							
Cannabis	20.68 (7.02, 34.33)	0.0030	-1541.24	1.9495	phi	0.3466	0.0002
Cigarettes: Alcohol	48.6 (2.75, 94.46)	0.0378			psi	0.1488	0.0006
Cigarettes: Cannabis	-46.18 (-84.76, -7.6)	0.0190			rho	-0.5248	2.4E-06
Alcohol	-25.69 (-44.01, -7.37)	0.0060			lambda	0.4837	1.3E-09
<i>4-Way Interactive model</i>							
<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine)</i>			phi		0.3169		0.0002
Cannabis	5.42 (2.34, 8.5)	0.0006	-1540.34	1.9470	psi	0.1479	0.0007
Alcohol	-8.18 (-14.61, -1.74)	0.0128			rho	-0.4896	3.1E-05
Cigarettes: Analgesics	-31.81 (-56.07, -7.54)	0.0102			lambda	0.4514	2.1E-07

<i>Interactive Full Model - 0 Lags</i>							
<i>spreml(Cancer Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6 Races+Income)</i>							
Cigarettes	28.41 (12.48, 44.34)	0.0005	-1520.55	1.8458	phi	0.1709	0.0017
Cannabis	45.67 (18.77, 72.56)	0.0009			psi	0.1079	0.0138
White	5.24 (3.38, 7.1)	0.0000			rho	-0.4106	0.0029
Cigarettes: Cannabis: Alcohol	840.86 (416.29, 1265.44)	0.0001			lambda	0.3643	0.0006
Alcohol: Analgesics	638.1 (283.09, 993.12)	0.0004					
Asian-American	0.6 (0.23, 0.97)	0.0015					
Hispanic-American	0.45 (0.11, 0.79)	0.0089					
Cigarettes: Cannabis: Analgesics	966.38 (184.69, 1748.06)	0.0154					
AIAN-American	-8.3 (-15.42, -1.18)	0.0224					
Cigarettes: Analgesics	-240.1 (-391.42, -88.77)	0.0019					
Cannabis: Alcohol: Analgesics	-2613.19 (-4248.03, -978.35)	0.0017					
Cigarettes: Cannabis	-235.06 (-381.19, -88.92)	0.0016					
Alcohol	-79.04 (-114.93, -43.14)	1.6E-05					

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Table 8.: Time-Lagged Spatiotemporal Models

Lagged Variables	Parameter			Model				
	Parameter	Estimate (C.I.)	P-Value	LogLik	S.D.	Model Parameter	Estimate	P-Value
	Full model - 2 Lags - Just Lagging Cannabis							
	<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6_Races+Income)</i>							
Cannabis, 2	Caucasian-American	5.3 (3.63, 6.97)	5.3E-10	-1329.42	1.8583	phi	0.1690	0.0037
	Asian-American	0.63 (0.31, 0.95)	1.3E-04			psi	0.1476	0.0018
	Hispanic-American	0.54 (0.21, 0.86)	0.0013			rho	-0.4435	8.3E-04
	AIAN-American	-11.33 (-18.34, -4.32)	0.0015			lambda	0.4234	9.1E-06
	Full model - 4 Lags - Just Lagging Cannabis							
	<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6_Races+Income)</i>							
Cannabis, 4	Caucasian-American	4.81 (2.92, 6.7)	6.1E-07	-1130.71	1.8616	phi	0.2095	0.0031
	Asian-American	0.67 (0.31, 1.03)	0.0003			psi	0.1134	0.0356
	Hispanic-American	0.55 (0.18, 0.92)	0.0037			rho	-0.5410	3.0E-05
	Cigarettes: Cannabis: Analgesics	261.1 (19.06, 503.15)	0.0345			lambda	0.4597	9.1E-07
	Cannabis: Analgesics	-107.34 (-193.09, -21.6)	0.0141					
	AIAN-American	-12.1 (-19.56, -4.64)	0.0015					
	Full model - 6 Lags - Just Lagging Cannabis							
	<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6_Races+Income)</i>							
Cannabis, 6	Caucasian-American	7.54 (3.96, 11.12)	3.6E-05	-936.96	1.9697	phi	0.2705	0.0022
	Asian-American	0.95 (0.34, 1.54)	0.0020			psi	0.0992	0.1012

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	Cannabis	8.49 (1.47, 15.5)	0.0177			rho	0.4222	0.0006
	Hispanic-American	0.7 (0.11, 1.29)	0.0202			lambda	-0.4083	0.0059
	Cannabis: Analgesics	-47.05 (-79.05, -15.03)	0.0040					
	Full Model - 1 Temporal Lag							
	spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6_Races+Income)							
Cigarettes, 1	Caucasian-American	5.42 (3.72, 7.12)	4.6E-10	-1426.33	1.8466	phi	0.1684	0.0027
Alcohol, 1	Asian-American	0.67 (0.33, 1)	0.0001			psi	0.1408	0.0016
Cannabis, 1	Hispanic-American	0.56 (0.22, 0.9)	0.0014			rho	-0.4380	0.0009
Analgesics, 1	Cannabis	7.88 (1.7, 14.06)	0.0125			lambda	0.4226	1.2E-05
Cocaine, 1	Cigarettes: Cannabis: Alcohol	182.23 (29.55, 334.9)	0.0193					
	AIAN-American	-9.03 (-16.1, -1.96)	0.0123					
	Cannabis: Alcohol	-114.28 (-198.3, -30.27)	0.0077					
	Full Model - 2 Temporal Lags							
	spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6_Races+Income)							
Cigarettes, 2	Caucasian-American	9.62 (6.82, 12.43)	1.7E-11	-1317.36	1.8519	phi	0.1408	0.0083
Alcohol, 2	Cigarettes: Cannabis: Alcohol	658.72 (396.6, 920.84)	8.4E-07			psi	0.1469	0.0018
Cannabis, 2	Asian-American	1.32 (0.75, 1.89)	6.4E-06			rho	0.3276	0.0126
Analgesics, 2	Alcohol: Analgesics	306.67 (143.27, 470.07)	0.0002			lambda	-0.2888	0.0462
Cocaine, 2	Hispanic-American	0.69 (0.26, 1.12)	0.0016					
	Income	-2.15 (-4.22, -0.08)	0.0415					
	Cannabis: Alcohol: Analgesics	-1810.02 (-2618.86, -1001.18)	1.2E-05					
	Cigarettes: Alcohol	-133.02 (-184.5, -81.54)	4.1E-07					
	Full Model - 4 Temporal Lags							
	spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6_Races+Income)							
Cigarettes, 4	Caucasian-American	5.25 (3.17, 7.33)	7.6E-07	-1129.73	1.8795	phi	0.1863	0.0058
Alcohol, 4	Cigarettes: Cannabis: Alcohol	472.69 (145.49, 799.88)	0.0046			psi	0.1341	0.0127

Cannabis, 4	Asian-American	0.56 (0.16, 0.95)	0.0055			rho	-0.4598	0.0040
Analgesics, 4	Hispanic-American	0.5 (0.13, 0.87)	0.0085			lambda	0.4021	8.6E-04
Cocaine, 4	Cigarettes: Alcohol: Analgesics	603.85 (143.88, 1063.82)	0.0101					
	Cigarettes: Alcohol	-80.89 (-138.89, -22.89)	0.0063					
	AIAN-American	-10.99 (-18.8, -3.18)	0.0058					
	Cigarettes: Cannabis: Alcohol: Analgesics	-3668.28 (-6170.15, -1166.42)	0.0041					
	Full Model - 6 Temporal Lags							
Cigarettes, 6	<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6 Races+Income)</i>							
Alcohol, 6	Caucasian-American	4.28 (2.17, 6.4)	7.4E-05	-938.093	1.9015	phi	0.2238	0.0053
Cannabis, 6	Asian-American	0.5 (0.13, 0.87)	0.0089			psi	0.1218	0.0448
Analgesics, 6	Hispanic-American	0.51 (0.12, 0.91)	0.0115			rho	-0.5495	7.7E-05
Cocaine, 6	AIAN-American	-11.64 (-19.61, -3.66)	0.0042			lambda	0.5042	1.8E-07

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1143**Table 9.: Spatially- and Temporally- Lagged Spatiotemporal Models**

Lagged Variables	Parameter			Model				
	Parameter	Estimate (C.I.)	P-Value	LogLik	S.D.	Model Parameter	Estimate	P-Value
	<i>Full Model - 1 Spatial & 1 Temporal Lag</i>							
	<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6 Races+Income)</i>							
Cigarettes, 1	Caucasian-American	4.49 (2.56, 6.41)	4.9E-06	-1422.64	1.8639	phi	0.1534	0.0041
Alcohol, 1	Hispanic-American	0.61 (0.26, 0.96)	0.0006			psi	0.1284	0.0043
Cannabis, Sp1	Cannabis: Analgesics	110.36 (37.53, 183.19)	0.0030			rho	-0.3379	0.0408
Analgesics, 1	Cigarettes: Cannabis: Alcohol	1688.83 (336.9, 3040.77)	0.0143			lambda	0.3229	0.0134
Cocaine, 1	Asian-American	0.46 (0.09, 0.83)	0.0146					
	Cannabis: Alcohol: Analgesics	-885.51 (-1625.8, -145.21)	0.0191					
	AIAN-American	-10.01 (-17.08, -2.94)	0.0055					
	Analgesics	-18.96 (-29.31, -8.61)	0.0003					
	<i>Full Model - 2 Spatial & 2 Temporal Lags</i>							
	<i>spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6 Races+Income)</i>							
Cigarettes, 2	Caucasian-American	8.03 (5.67, 10.39)	2.6E-11	-1319.97	1.8579	phi	0.0990	0.0324
Alcohol, 2	Asian-American	1.02 (0.53, 1.51)	5.2E-05			psi	0.1426	0.0032
Cannabis, Sp2	Hispanic-American	0.66 (0.33, 0.99)	9.0E-05			rho	-0.2287	0.3086
Analgesics, 2	Analgesics	55.5 (26.18, 84.81)	0.0002			lambda	0.2307	0.2080
Cocaine, 2	Cigarettes: Cannabis: Alcohol: Analgesics	3954.04 (1565.01, 6343.08)	0.0012					
	Cocaine	15.51 (1.58, 29.44)	0.0291					
	Cigarettes: Cannabis: Analgesics	-749.24 (-1219.42, -279.07)	0.0018					

	Alcohol: Analgesics	-377.69 (-553.03, -202.35)	2.4E-05					
	Full Model - 4 Spatial & Temporal Lags							
	spreml(Cancer_Rate ~ Cigarettes * Cannabis * Alcohol * Analgesics + Cocaine+ 6 Races+Income)							
Cigarettes, 4	Caucasian-American	5.18 (3.29, 7.07)	7.7E-08	-1133.35	1.8790	phi	0.1850	0.0049
Alcohol, 4	Asian-American	0.59 (0.25, 0.93)	0.0008			psi	0.1286	0.0176
Cannabis, Sp4	Hispanic-American	0.52 (0.16, 0.87)	0.0045			rho	-0.4868	0.0004
Analgesics, 4	Alcohol: Analgesics	-27.25 (-54.07, -0.43)	0.0464			lambda	0.4290	2.5E-05
Cocaine, 4	AIAN-American	-10.96 (-18.4, -3.51)	0.0039					

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1152**Table 10.: Spatially- and Temporally- Lagged Spatiotemporal Models**

Lagged Variables	Parameter			Model				
	Parameter	Estimate (C.I.)	P-Value	LogLik	S.D.	Model Parameter	Estimate	P-Value
	<i>Cannabinoids</i>							
	<i>Cannabinoids as Main Effects</i>							
	<i>spreml(Cancer Rate ~ Cigarettes * THC * Cannabigerol * Alcohol + Analgesics + Cocaine)</i>							
	Caucasian-American	4.83 (2.77, 6.89)	4.5E-06	-1511.96	1.8350	phi	0.2050	0.0009
	Cigarettes: Alcohol	334 (171.12, 496.88)	0.0001			psi	0.0889	0.0450
	Alcohol: Analgesics	312 (149.91, 474.09)	0.0002			rho	-0.4495	0.0003
	Cigarettes: Δ9THC: Analgesics	391 (181.28, 600.72)	0.0003			lambda	0.3639	0.0001
	Δ9THC: Alcohol	116 (51.71, 180.29)	0.0004					
	Cigarettes: Δ9THC: Cannabigerol: Alcohol	4810 (2124.8, 7495.2)	0.0004					
	Δ9THC: Cannabigerol	109 (41.58, 176.42)	0.0016					
	Analgesics	96.5 (35.74, 157.26)	0.0018					
	Asian-American	0.57 (0.19, 0.94)	0.0029					
	Δ9THC: Cannabigerol: Alcohol: Analgesics	5640 (1680.8, 9599.2)	0.0052					
	Hispanic-American	0.41 (0.07, 0.76)	0.0193					
	Cigarettes: Δ9THC	9.01 (0.6, 17.42)	0.0359					
	AIAN-American	-8.84 (-16.13, -1.55)	0.0175					
	Cigarettes: Δ9THC: Cannabigerol: Alcohol: Analgesics	-18100 (-29977.6, -6222.4)	0.0028					
	Cigarettes: Δ9THC: Cannabigerol	-385 (-612.36, -157.64)	0.0009					
	Δ9THC: Cannabigerol: Alcohol	-1480 (-2346.32, -613.68)	0.0008					
	Δ9THC: Analgesics	-130 (-199.78, -60.22)	0.0003					
	Cigarettes: Δ9THC: Alcohol	-384 (-583.92, -184.08)	0.0002					
	Cigarettes: Analgesics	-383 (-563.32, -202.68)	3.2E-05					

	Alcohol	-137 (-197.96, -76.04)	1.1E-05					
<i>Cannabinoids as Main Effects - 2 Lags</i>								
<i>spreml(Cancer_Rate ~ Cigarettes * THC * Cannabigerol * Alcohol + Analgesics + Cocaine)</i>								
THC, 2	Caucasian-American	4.63 (2.53, 6.72)	1.5E-05	-1320.47	1.8880	phi	0.1976	0.0021
Cannabigerol, 2	Cannabigerol	21.6 (9.29, 33.9)	0.0006			psi	0.1322	0.0052
	THC: Alcohol	10.25 (4.12, 16.37)	0.0010			rho	-0.3332	0.0881
	Asian-American	0.56 (0.17, 0.95)	0.0053			lambda	0.3037	0.0500
	Cannabigerol: Alcohol: Analgesics	1176.24 (308.66, 2043.82)	0.0079					
	Hispanic-American	0.47 (0.1, 0.84)	0.0117					
	AIAN-American	-10.84 (-18.51, -3.17)	0.0056					
	Cannabigerol: Alcohol	-288.07 (-474.04, -102.11)	0.0024					
	THC: Cannabigerol: Analgesics	92.38 (39.25, 145.5)	0.0007					
<i>Cannabinoids as Main Effects - 4 Lags</i>								
<i>spreml(Cancer_Rate ~ Cigarettes * THC * Cannabigerol * Alcohol + Analgesics + Cocaine)</i>								
THC, 4	Caucasian-American	4.31 (2.26, 6.36)	3.9E-05	-1126.72	1.8642	phi	0.1876	0.0047
Cannabigerol, 4	Cigarettes: THC	2.87 (1.47, 4.27)	5.9E-05			psi	0.1223	0.0246
	Asian-American	0.64 (0.26, 1.03)	0.0010			rho	-0.4917	0.0007
	Hispanic-American	0.58 (0.21, 0.95)	0.0021			lambda	0.3940	0.0004
	Cigarettes: Cannabigerol: Alcohol	668.38 (191.09, 1145.67)	0.0061					
	Cigarettes	-3.45 (-5.85, -1.04)	0.0050					
	AIAN-American	-11.66 (-19.15, -4.17)	0.0023					
	Cannabigerol: Alcohol	-329.3 (-523.66, -134.95)	0.0009					
<i>Cannabinoids as Main Effects - 6 Lags</i>								
<i>spreml(Cancer_Rate ~ Cigarettes * THC * Cannabigerol * Alcohol + Analgesics + Cocaine)</i>								
THC, 6	Cigarettes: THC	28.16 (18.61, 37.71)	7.6E-09	-918.382	1.8922	phi	0.2868	0.0023
Cannabigerol, 6	THC: Cannabigerol	46.22 (30.06, 62.38)	2.1E-08			psi	0.1197	0.0495

	Asian-American	0.67 (0.21, 1.12)	0.0039			rho	-0.5066	0.0004
	Caucasian-American	3.19 (1.01, 5.38)	0.0042			lambda	0.3707	0.0007
	Cocaine	18.22 (3.99, 32.46)	0.0121					
	Cigarettes: Cannabigerol: Alcohol	724.22 (143.74, 1304.71)	0.0145					
	AIAN-American	-10.25 (-18.78, -1.73)	0.0184					
	Cannabigerol: Alcohol	-329.39 (-580.71, -78.07)	0.0102					
	Cigarettes: THC: Cannabigerol	-177.1897 (-248.1, -106.28)	9.7E-07					
	THC	-7.21 (-9.86, -4.55)	1.1E-07					
	Cigarettes	-29.01 (-39.21, -18.82)	2.4E-08					

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Table 12.: Spatially- and Temporally- Lagged Spatiotemporal Models

Parameter	Estimate (C.I.)	R.R. (C.I.)	E-Values
<i>LINEAR REGRESSION</i>			
<i>Cancer Rate Over Time</i>			
Year	0.14 (0.1, 0.17)	1.06 (1.04, 1.08)	1.31, 1.27
<i>Cancer Rate by Δ9THC</i>			
Δ 9THC	0.33 (0.15, 0.5)	1.15 (1.07, 1.23)	1.55, 1.33
<i>Cancer Rate by Drug Rate</i>			
Drug_Rate: Cannabis	4.63 (2.11, 7.15)	6.83 (2.41, 19.41)	13.15, 4.25
<i>Cancer Rate by Cannabinoid Over Time</i>			
Cannabinol	6.54 (5.07, 8.01)	15.54 (8.39, 28.78)	30.58, 16.27
Cannabigerol	7.65 (5.91, 9.38)	24.71 (11.96, 51.02)	48.91, 23.41
Drug_Rate	2.14 (1.55, 2.73)	2.45 (1.91, 3.14)	4.34, 3.24
Cannabichromene	3.86 (0.29, 7.42)	5.04 (1.14, 22.44)	9.54, 1.51
<i>Cancer Rate by Ethnic Cannabis Exposure</i>			
Ethnic_THC_Exposure	0.14 (0.07, 0.21)	1.06 (1.03, 1.09)	1.31, 1.20
Asian-Am_THC_Exposure	0.28 (0.02, 0.55)	1.12 (1.01, 1.26)	1.50, 1.10
<i>Legal Status</i>			
Decriminalized	0.85 (0.44, 1.26)	1.42 (1.20, 1.69)	2..20, 1.69
Liberal	0.663 (0.35, 0.98)	1.32 (1.15, 1.50)	1.96, 1.58
Legal	1.3286 (0.47, 2.19)	1.73 (1.21, 2.45)	2.86, 1.72
<i>Cancer by Legal Status</i>			
Decriminalized	0.78 (0.37, 1.19)	1.38 (1.16, 1.64)	2.11, 1.60
Legal	1.51 (0.68, 2.35)	1.87 (1.33, 2.66)	3.16, 1.98
<i>Cancer by Year * Status</i>			

Year: Decriminalized	0.0003 (0.0001, 0.0005)	1.00013 (1.00004, 1.00021)	1.011, 1.006
<i>Cancer by Year * Dichotomized Status</i>			
Year: Liberal	0.0002 (0, 0.0004)	1.00008 (1.00001, 1.00015)	1.0090, 1.0035
<i>MIXED EFFECTS REGRESSION</i>			
<i>Additive Model</i>			
Cannabis	5.34 (0.07, 10.6)	4.11 (1.02, 16.59)	7.70, 1.18
<i>Interactive Drugs Model</i>			
Cannabis	72.88 (49.6, 96.15)	5.02E+08 (8.45E+05, 2.97E+11)	1.01E+09, 1.69E+06
Cigarettes: Cannabis: Analgesics	2788.19 (1676.17, 3900.2)	Infinity (2.40E+200, Infinity)	Infinity, Infinity
<i>Full Interactive Model</i>			
Cannabis	79.27 (56.77, 101.78)	7.00E+09 (1.14E+07, 4.31E+12)	1.40E+09, 2.27E+07
Cigarettes: Cannabis: Analgesics	2525.7 (1488.65, 3562.75)	Infinity (1.38E+185, Infinity)	Infinity, Infinity
<i>Full Interactive Model with Cannabinoids</i>			
Cigarettes: Cannabigerol: Alcohol	4741.19 (3077.86, 6404.51)	Infinity (Infinity, Infinity)	Infinity, Infinity
Δ9THC: Alcohol	14.95 (7.74, 22.16)	73.78 (9.31, 584.34)	147.07, 18.12
Cigarettes: Δ9THC	26.57 (15.54, 37.6)	2.09E+03 (87.97, 4.95E+04)	4.18E+03, 175.45
<i>GEOSPATIAL REGRESSION</i>			
<i>Additive Model</i>			
Cannabis	5.16 (2.26, 8.06)	11.18 (2.89, 43.30)	21.84, 5.22
<i>3-Way Interactive model</i>			
Cannabis	20.68 (7.02, 34.33)	1.55E+04 (26.85, 9.01E+06)	3.11E+04, 53.19
<i>4-Way Interactive model</i>			
Cannabis	5.42 (2.34, 8.5)	12.61 (2.99, 53.07)	24.71, 5.45
<i>Interactive Full Model - 0 Lags</i>			
Cannabis	45.67 (18.77, 72.56)	6.00E+10 (1.07E+04, 3.45E+15)	1.20E+120, 5.15E+04
Cigarettes: Cannabis: Alcohol	840.86 (416.29, 1265.44)	1.09E+180 (2.07E+89, 5.78E+270)	Infinity, 4.14E+89
Cigarettes: Cannabis: Analgesics	966.38 (184.69, 1748.06)	8.18E+206 (7.64E+39, Infinity)	Infinity, 1.52E+40

Time Lagged Models			
Full model - 4 Lags - Just Lagging Cannabis			
Cigarettes: Cannabis: Analgesics	261.1 (19.06, 503.15)	8.26E+39 (0.07, 9.56E+80)	1.65E+40, 1.00
Full model - 6 Lags - Just Lagging Cannabis			
Cannabis	8.49 (1.47, 15.5)	50.45 (1.98, 1.28E+03)	100.41, 3.39
Full Model - 1 Temporal Lag			
Cannabis	7.88 (1.7, 14.06)	48.60 (2.32, 1.016E+03)	96.68, 4.07
Cigarettes: Cannabis: Alcohol	182.23 (29.55, 334.9)	1.00E+39 (2.45E+06, 4.07E+71)	1.99E+39, 4.91E+06
Full Model - 2 Temporal Lags			
Cigarettes: Cannabis: Alcohol	658.72 (396.6, 920.84)	3.76E+140 (5.65E+84, 2.53E+196)	7.58E+140, 1.13E+85
Full Model - 4 Temporal Lags			
Cigarettes: Cannabis: Alcohol	472.69 (145.49, 799.88)	9.42E+126 (2.81E+30, 3.15E+223)	1.88E+127, 5.62E+30
Space-Time Lagged Models			
Full Model - 1 Spatial & 1 Temporal Lag			
Cannabis: Analgesics	110.36 (37.53, 183.19)	2.51E+23 (9.78E+07, 6.48E+38)	5.03E+23, 1.95E+08
Cigarettes: Cannabis: Alcohol	1688.83 (336.9, 3040.77)	Infinity (1.033E+72, Infinity)	Infinity, 2.07E+72
Full Model - 2 Spatial & 2 Temporal Lags			
Cigarettes: Cannabis: Alcohol: Analgesics	3954.04 (1565.01, 6343.08)	Infinity (Infinity, Infinity)	Infinity, Infinity
Cannabinoid Models			
Cannabinoids as Main Effects			
Cigarettes: Δ9THC: Analgesics	391 (181.28, 600.72)	1.62E+84 (1.65E+39, 1.59E+129)	3.24E+84, 3.30E+39
Δ9THC: Alcohol	116 (51.71, 180.29)	1.11E+25 (1.68E+11, 7.41E+38)	2.23E+25, 3.36E+11
Cigarettes: Δ9THC: Cannabigerol: Alcohol	4810 (2124.8, 7495.2)	Infinity (Infinity, Infinity)	Infinity, Infinity
Δ9THC: Cannabigerol	109 (41.58, 176.42)	2.45E+23 (7.67E+08, 7.83E+37)	4.90E+23, 1.54E+09
Δ9THC: Cannabigerol: Alcohol: Analgesics	5640 (1680.8, 9599.2)	Infinity (Infinity, Infinity)	Infinity, Infinity
Cigarettes: Δ9THC	9.01 (0.6, 17.42)	87.15 (1.35, 5.61E+03)	173.80, 2.04

<i>Cannabinoids as Main Effects - 2 Lags</i>			
Cannabigerol	21.6 (9.29, 33.9)	3.32E+04 (89.18, 1.23E+07)	6.64E+04, 177.84
THC: Alcohol	10.25 (4.12, 16.37)	139.58 (7.34, 2.65E+03)	278.66, 14.15
Cannabigerol: Alcohol: Analgesics	1176.24 (308.66, 2043.82)	1.66E+246 (9.51E+64, Infinity)	Infinity, 1.91E+65
<i>Cannabinoids as Main Effects - 4 Lags</i>			
Cigarettes: THC	2.87 (1.47, 4.27)	4.06 (2.06, 8.04)	7.58, 3.52
Cigarettes: Cannabigerol: Alcohol	668.38 (191.09, 1145.67)	5.01E+141 (5.21E+40, 4.82E+242)	1.00E+142, 1.04E+41
<i>Cannabinoids as Main Effects - 6 Lags</i>			
Cigarettes: THC	28.16 (18.61, 37.71)	7.61E+05 (7.76E+03, 7.46E+07)	1.52E+06, 1.55E+04
THC: Cannabigerol	46.22 (30.06, 62.38)	4.50E+09 (1.92E+06, 1.06E+13)	9.01E+09, 3.84E+06
Cigarettes: Cannabigerol: Alcohol	724.22 (143.74, 1304.71)	1.82E+151 (1.84E+30, 1.80E+272)	3.64E+151, 3.68E+30
<i>Ethnicity Models</i>			
<i>Cancer Incidence as a Function of Racial Cannabis Exposure</i>			
Afric-Am. THC Exp: Hispan.Am THC Exp	1.74 (1.18, 2.29)	2.22 (1.72, 2.86)	3.86, 2.86
Afric-Am. THC Exp: Hispan.Am THC Exp: Asian-Am. THC Exp: AIAN-Am. THC Exp	0.15 (0.09, 0.21)	1.51 (1.18, 1.91)	2.38, 1.66
Asian-Am. THC Exp: AIAN-Am. THC Exp	0.89 (0.37, 1.41)	1.06 (1.04, 1.10)	1.34, 1.24

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Table 13.: Linear Regressions for Legal Status

Parameter Estimates			Model Parameters			
Parameter	Estimate (C.I.)	P-Value	R-Squared	F	dF	P-Value
<i>Cancer by Status</i>						
<i>lm(Cancer_Rate ~ Legal_Status)</i>						
Decriminalized	0.78 (0.37, 1.19)	2.0E-04	0.0268	7.88	3,746	3.49E-05
Legal	1.51 (0.68, 2.35)	4.0E-04				
<i>Cancer by Year * Status</i>						
<i>lm(Cancer_Rate ~ Year * Legal_Status)</i>						
Year	0.13 (0.09, 0.16)	4.3E-11	0.0809	17.5	4,745	1.01E-13
Year: Decriminalized	0.0003 (0.0001, 0.0005)	4.4E-03				
<i>Cancer by Year * Dichotomized_Status</i>						
<i>lm(Cancer_Rate ~ Year * Dichotomized_Status)</i>						
Year	0.128 (0.09, 0.16)	9.8E-12	0.0778	32.6358	2,747	2.58E-14
Year: Liberal	0.0002 (0, 0.0004)	2.1E-02				

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Figure Captions

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1175 Figure 1.: Pediatric Cancers 1975-2017, CDC SEER Explorer Dataset, USA National Level,

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data derived from 9 cancer registries.

Figure 2.: Drug use over time. Data from NSDUH 2002-2017, SAMHSA.

Figure 3.: Cannabinoid concentrations in Federal Seizures of Cannabis over time, Drug Enforcement Agency data [47-49].

Figure 4.: Total pediatric cancer incidence rate as a function of drug exposure.

Figure 5.: Total pediatric cancer incidence rate as a function of estimated state level cannabinoid exposure.

Figure 6.: Total pediatric cancer incidence rate as a function of estimated ethnic THC exposure.

Figure 7.: Total pediatric cancer incidence rate by cannabis use quintiles. (A) Boxplot over aggregated time. (B) Scatterplot over time by cannabis use quintiles. (C) Boxplot by dichotomized cannabis use quintiles, highest two quintiles vs. the lowest three. Note non-overlapping notches indicating significant differences. (D) Scatterplot over time of total pediatric cancer incidence rate by dichotomized cannabis use quintiles.

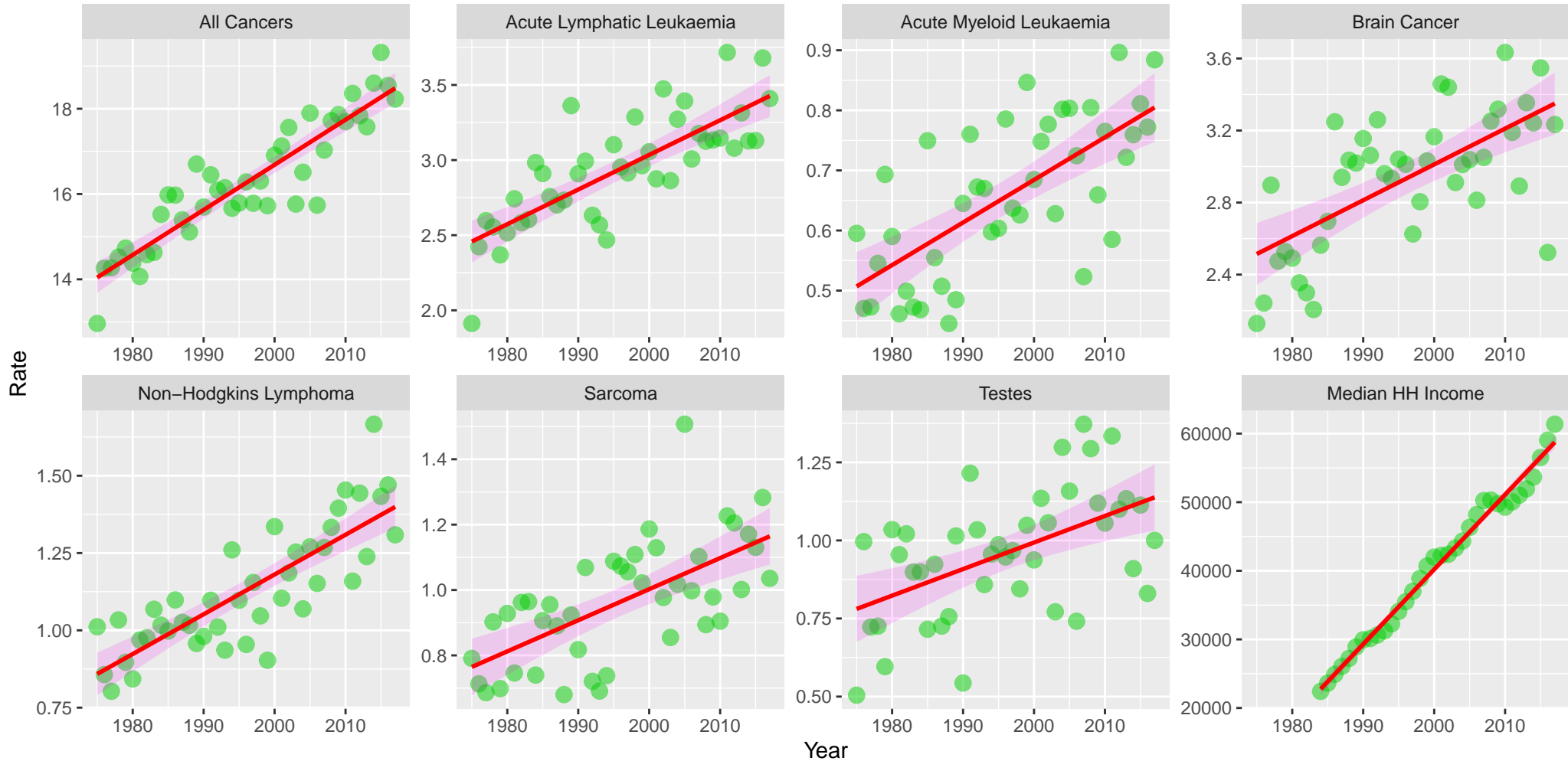
Figure 8.: Map graph of total pediatric cancer incidence rate by state over time sequence, by year.

Figure 9.: Geospatial linkages used for geospatiotemporal regression analyses. Note Alaska and Hawaii elided arithmetically onto continental USA. (A) Edited spatial links. (B) Final links.

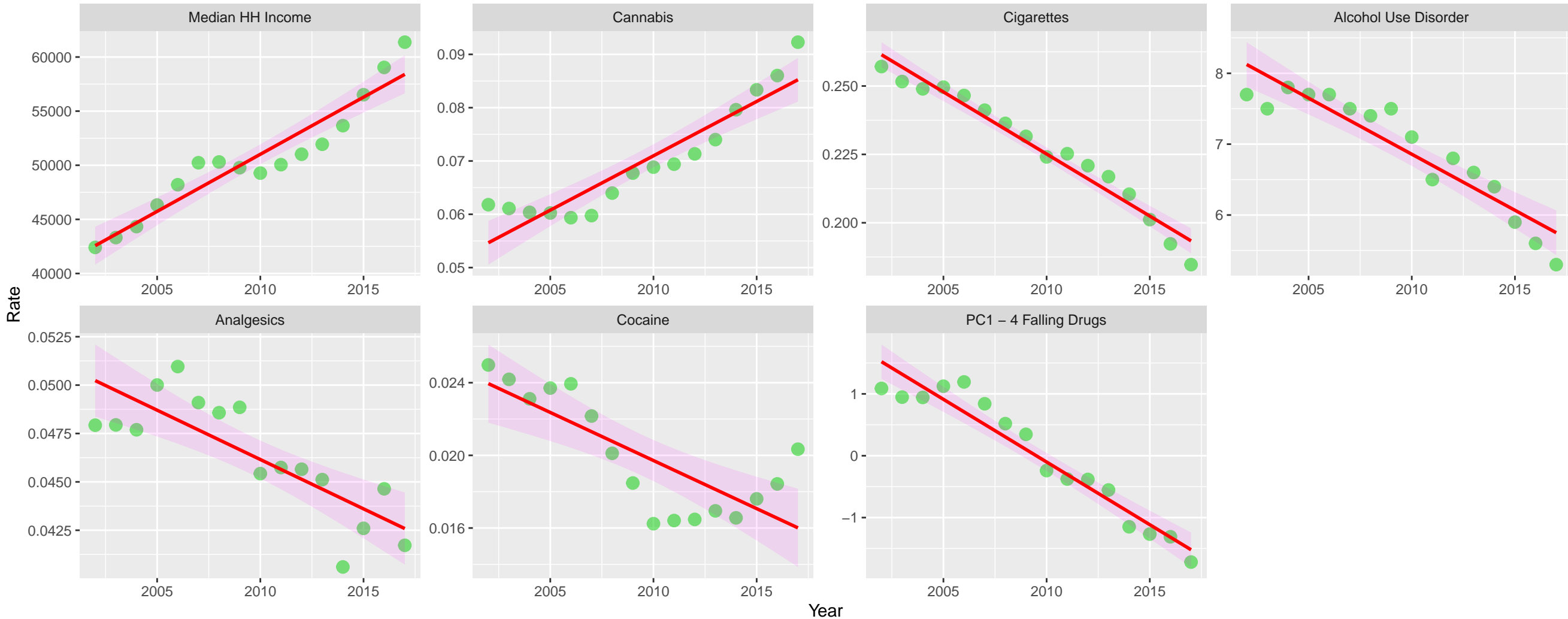
Figure 10: Effect of Cannabis Legal Status on total pediatric cancer incidence rate. (A) Scatterplot of legal statuses over time. (B) Scatterplot of legal status over time dichotomized as illegal status vs. liberal regimes.

Selected and Major Paediatric Cancer Rates 1975–2000

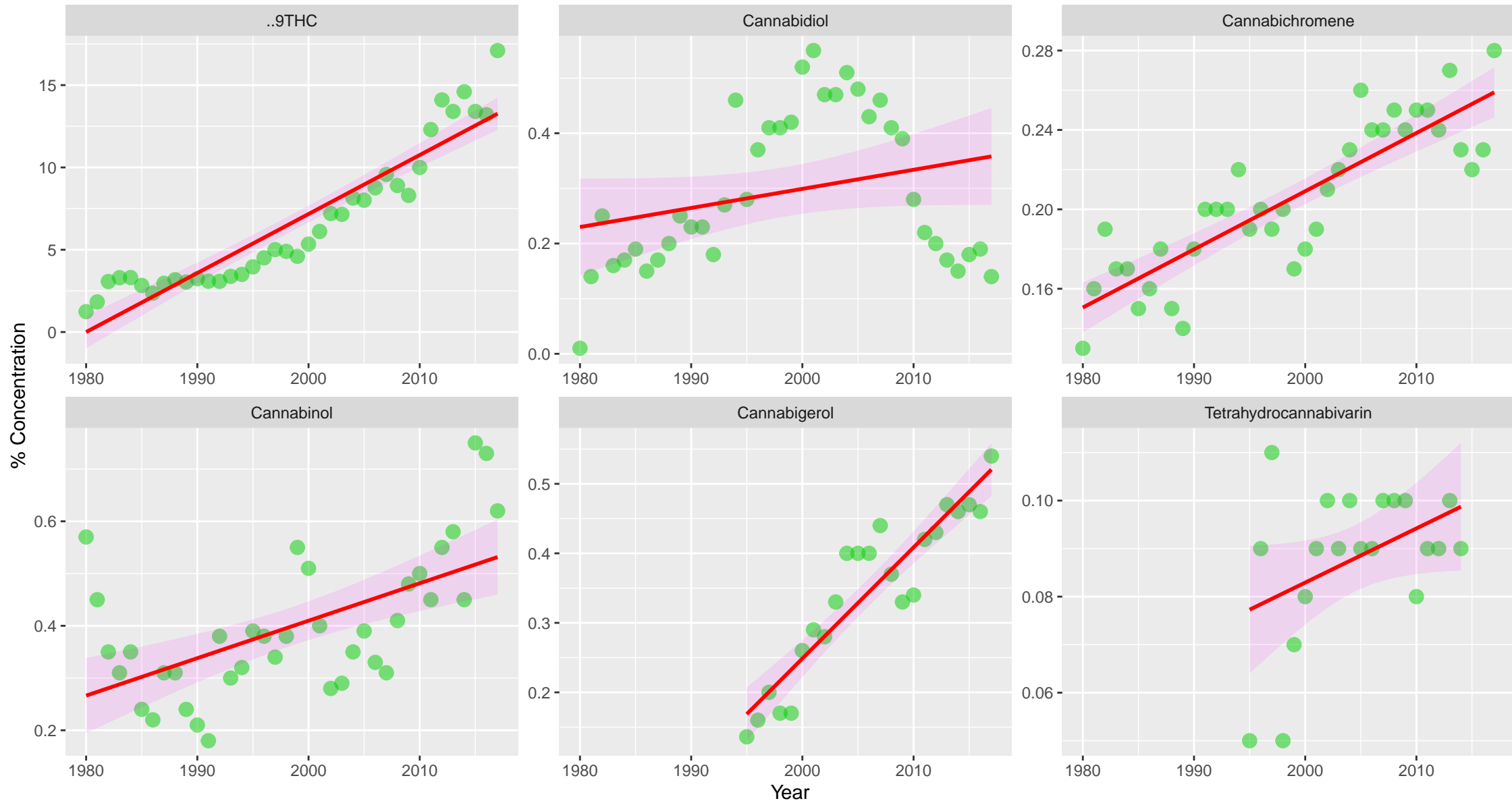
9 US Cancer Registries



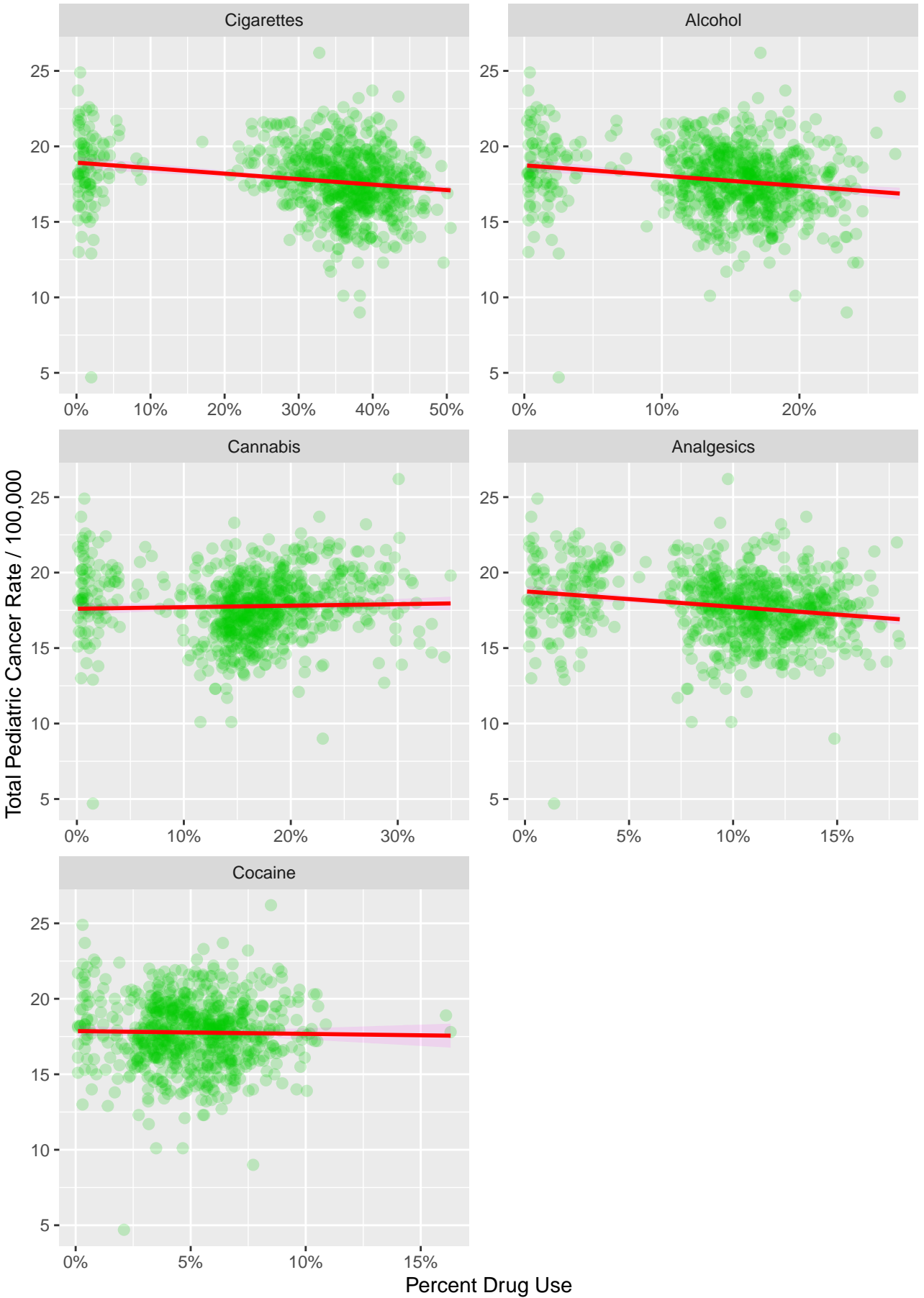
Drug Use and Income Over Time



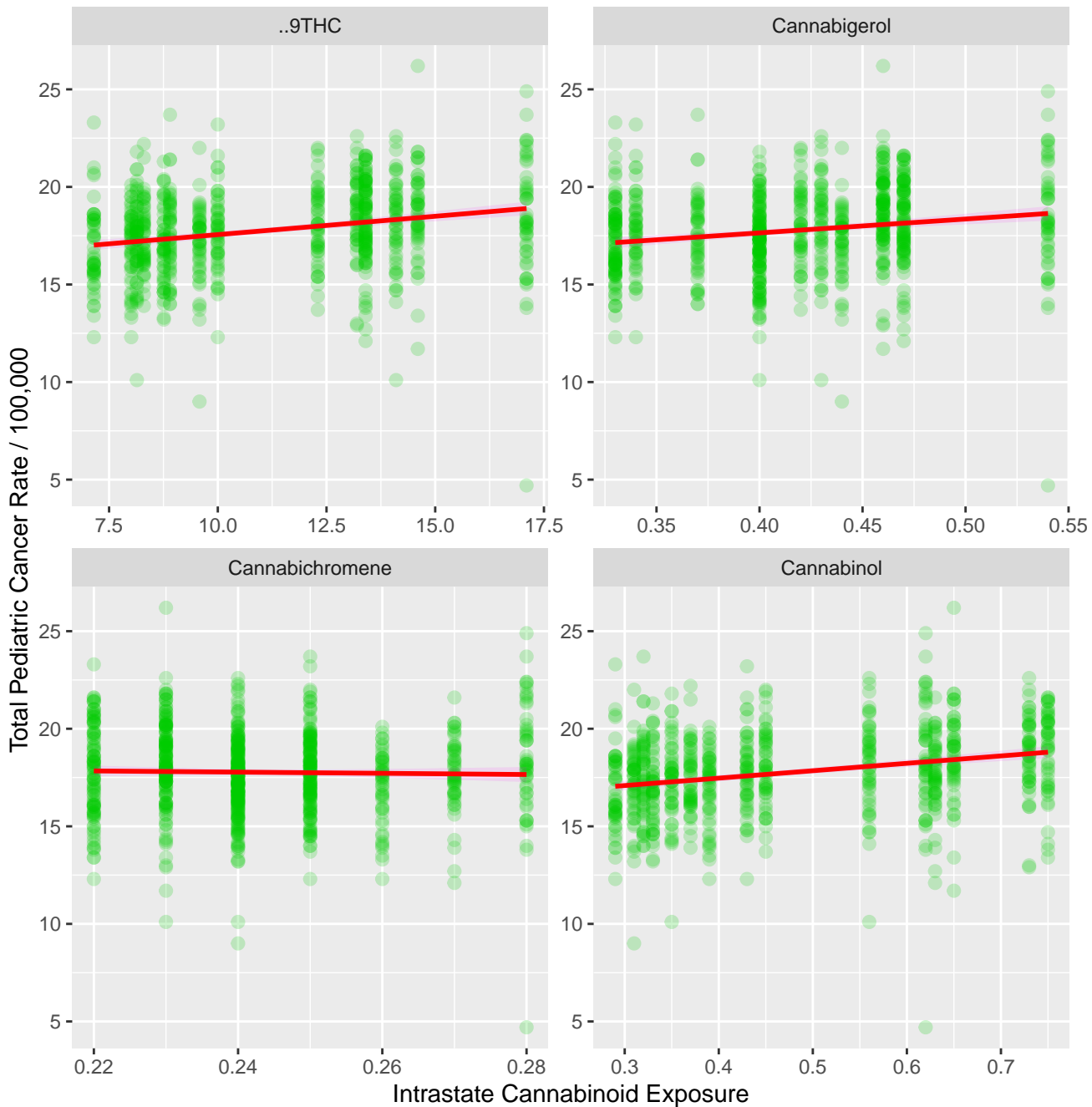
US National Cannabinoid Concentration Trends DEA Seizure Analyses



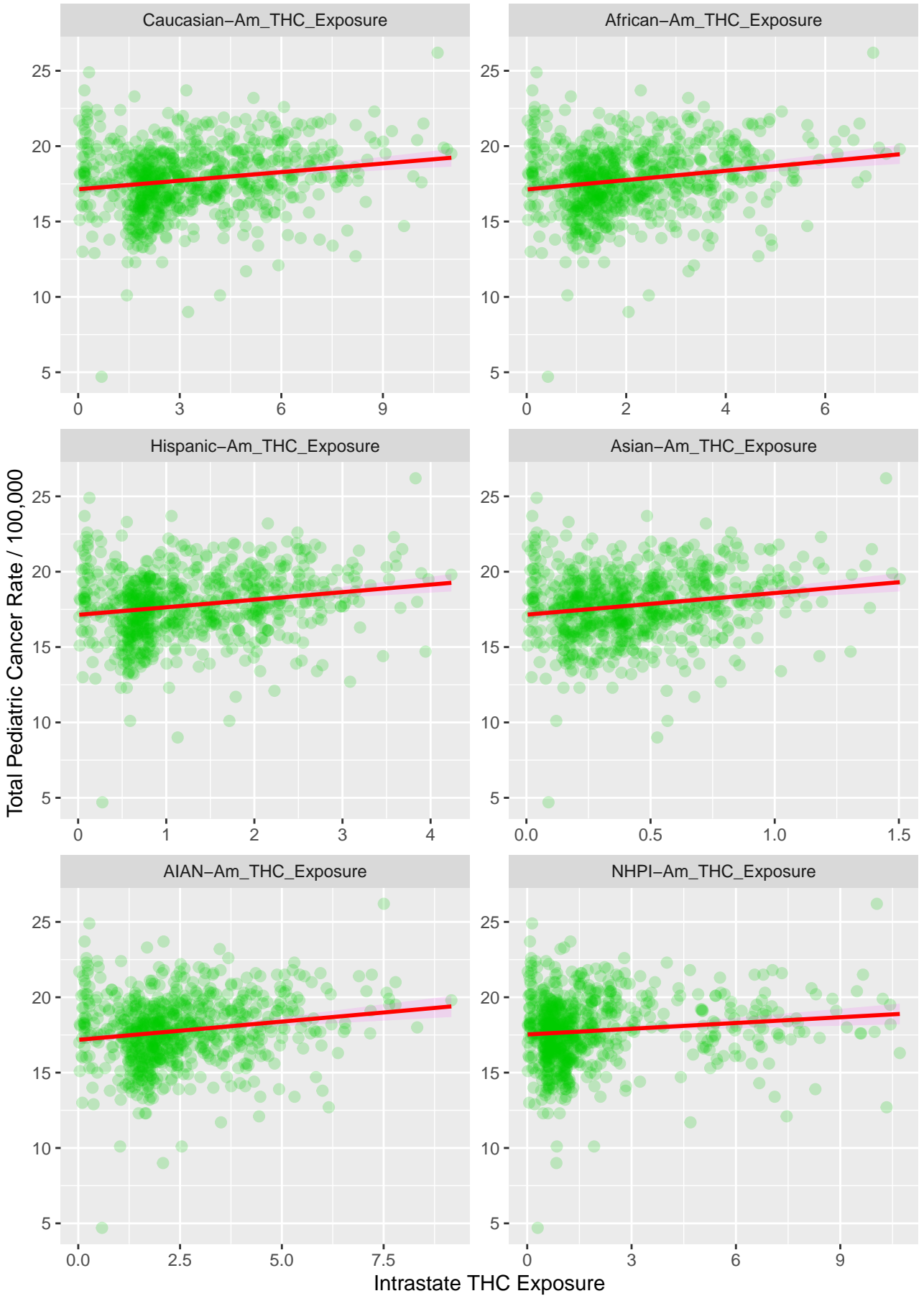
Pediatric Cancer Rate by Substance Exposure



Pediatric Cancer Rate by Cannabinoid Exposure

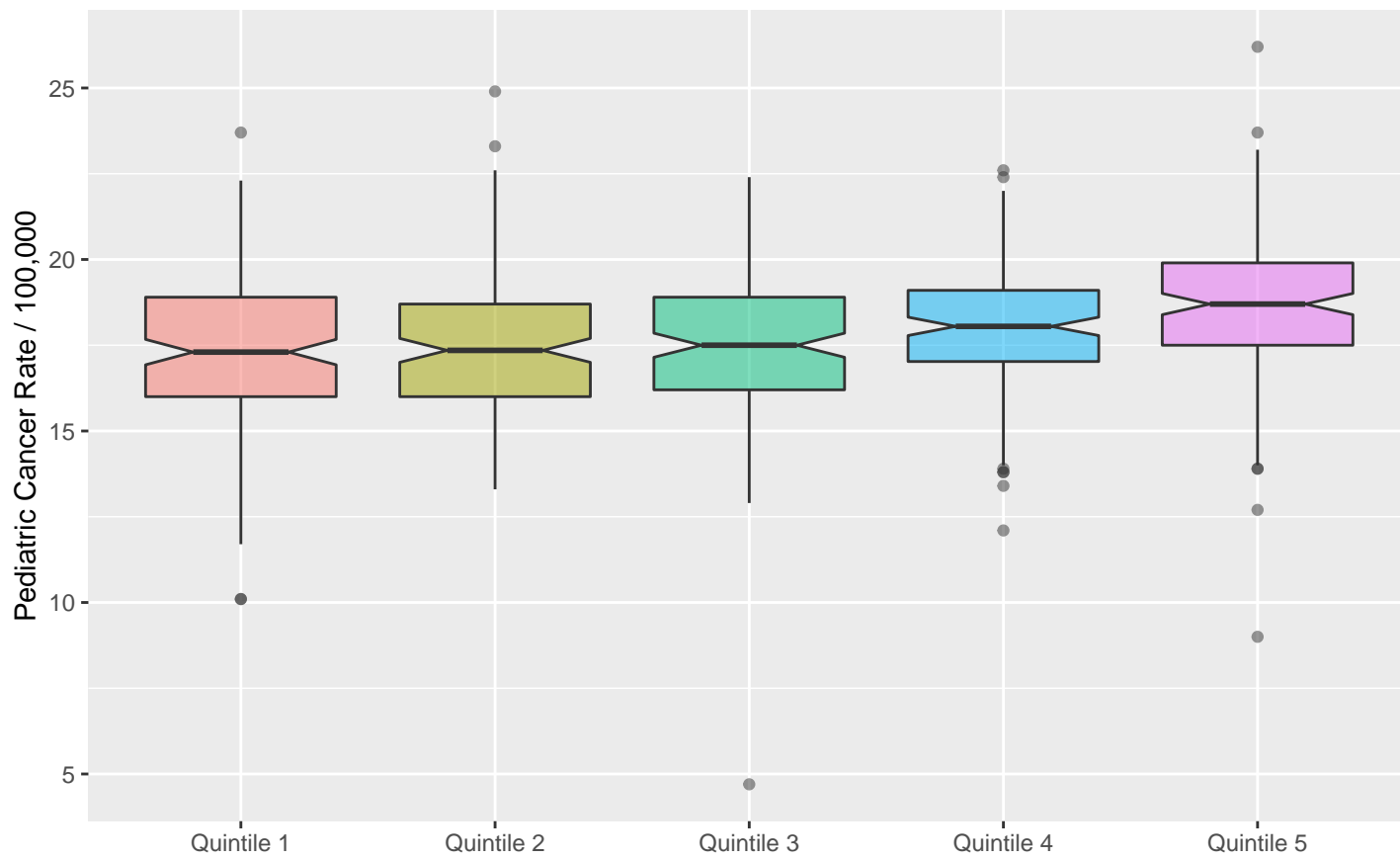


Pediatric Cancer Rate by Ethnic THC Exposure

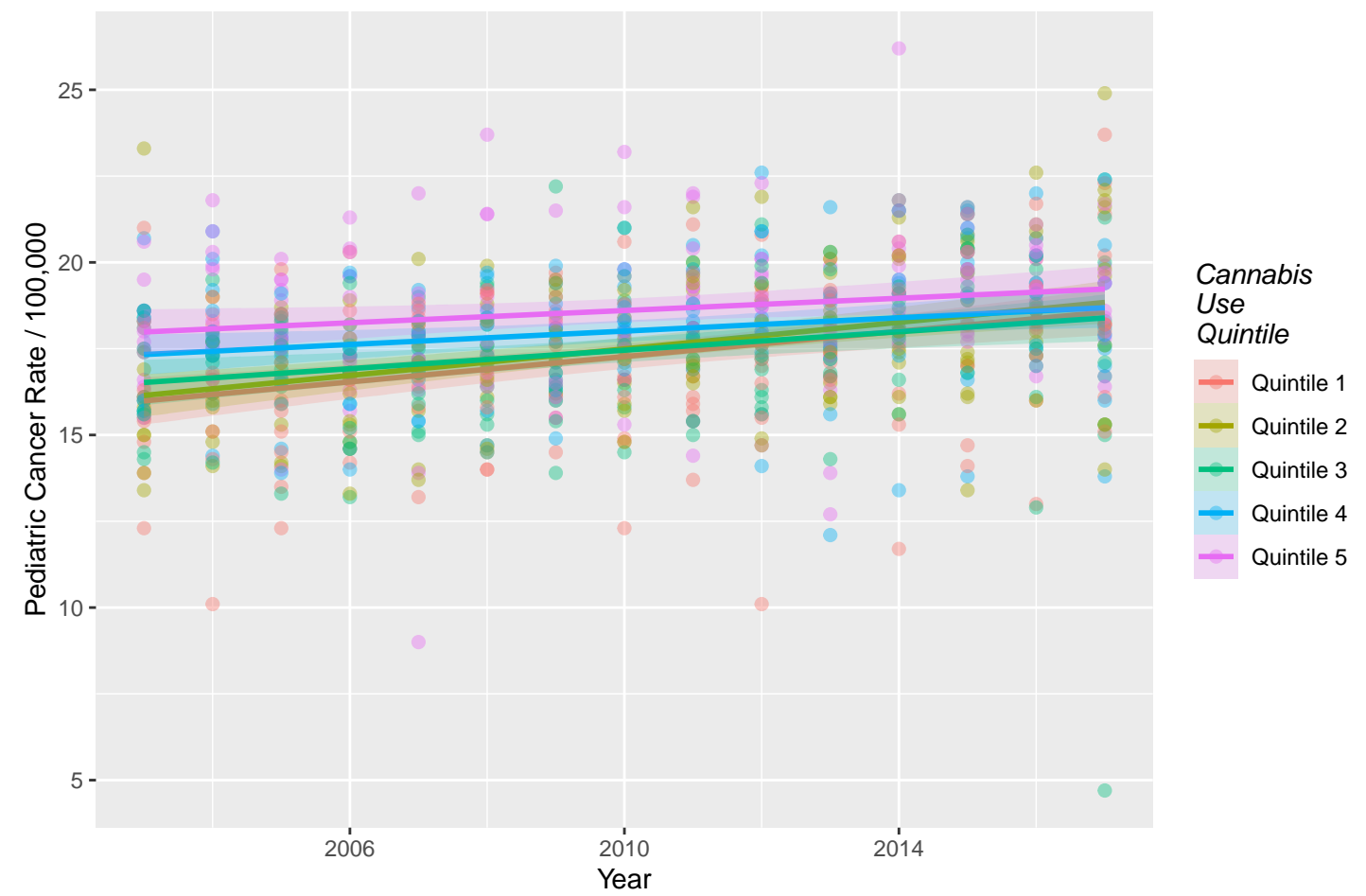


Pediatric Cancer Rate by Quintiles of Cannabis Use

Data – NCI SEER Program and NSDUH, SAMHSA

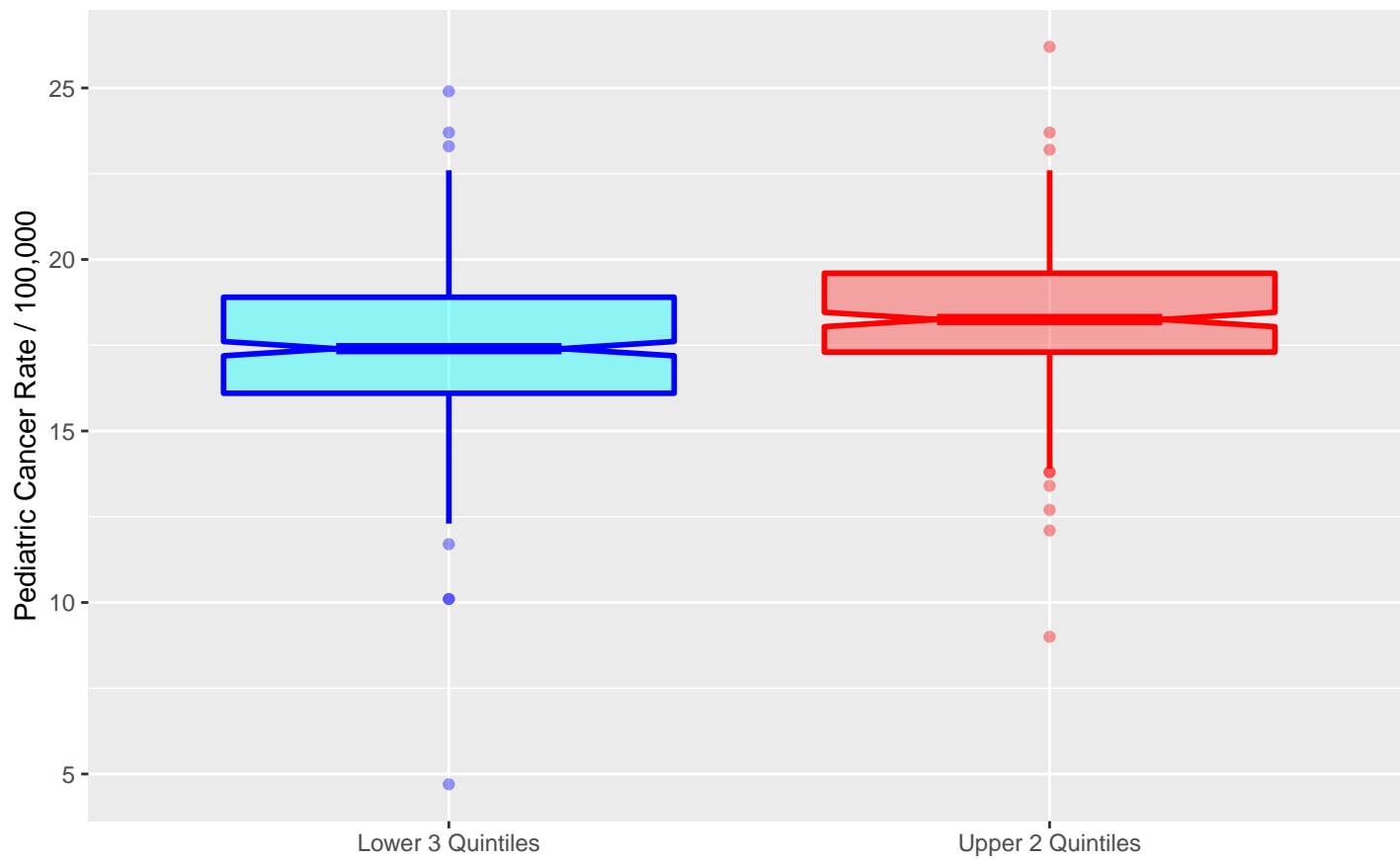


US Pediatric Cancer Rates by Cannabis Use Quintile

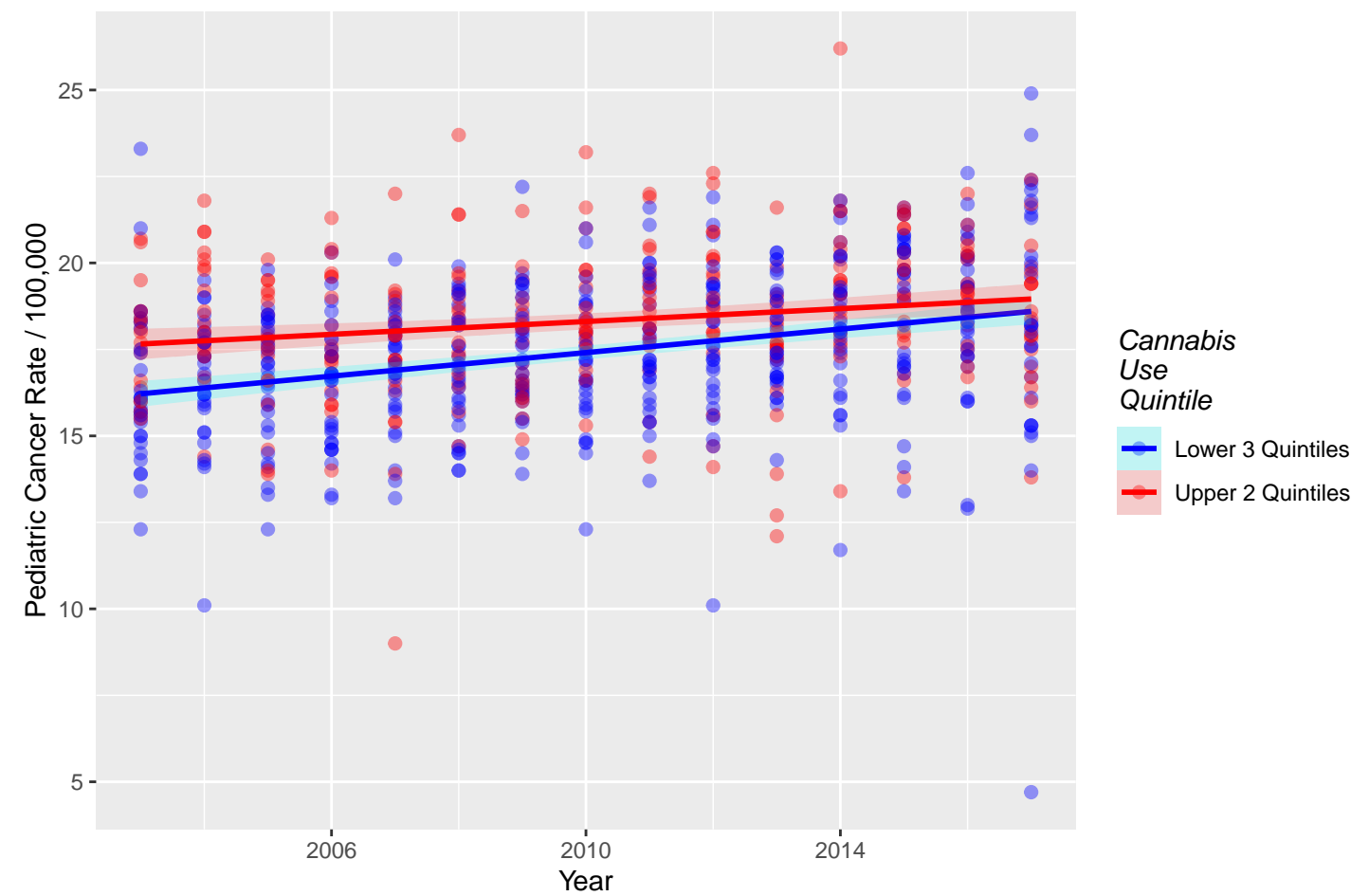


Pediatric Cancer Rate by Quintiles of Cannabis Use

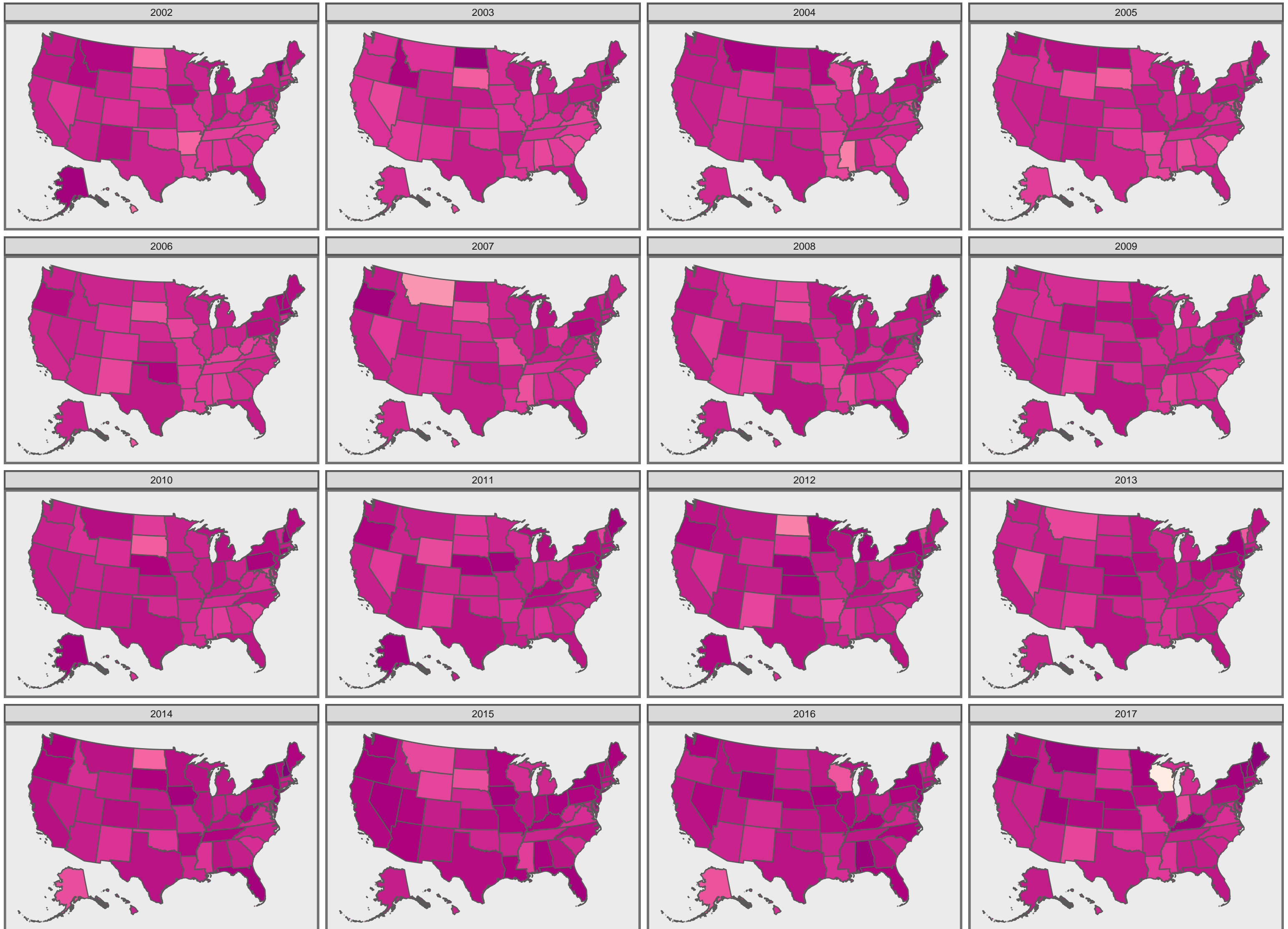
Data – NCI SEER Program and NSDUH, SAMHSA



US Pediatric Cancer Rates by Cannabis Use Quintile – Dichotomized



Log (Rates) of Paediatric Cancer US States 2002–2017



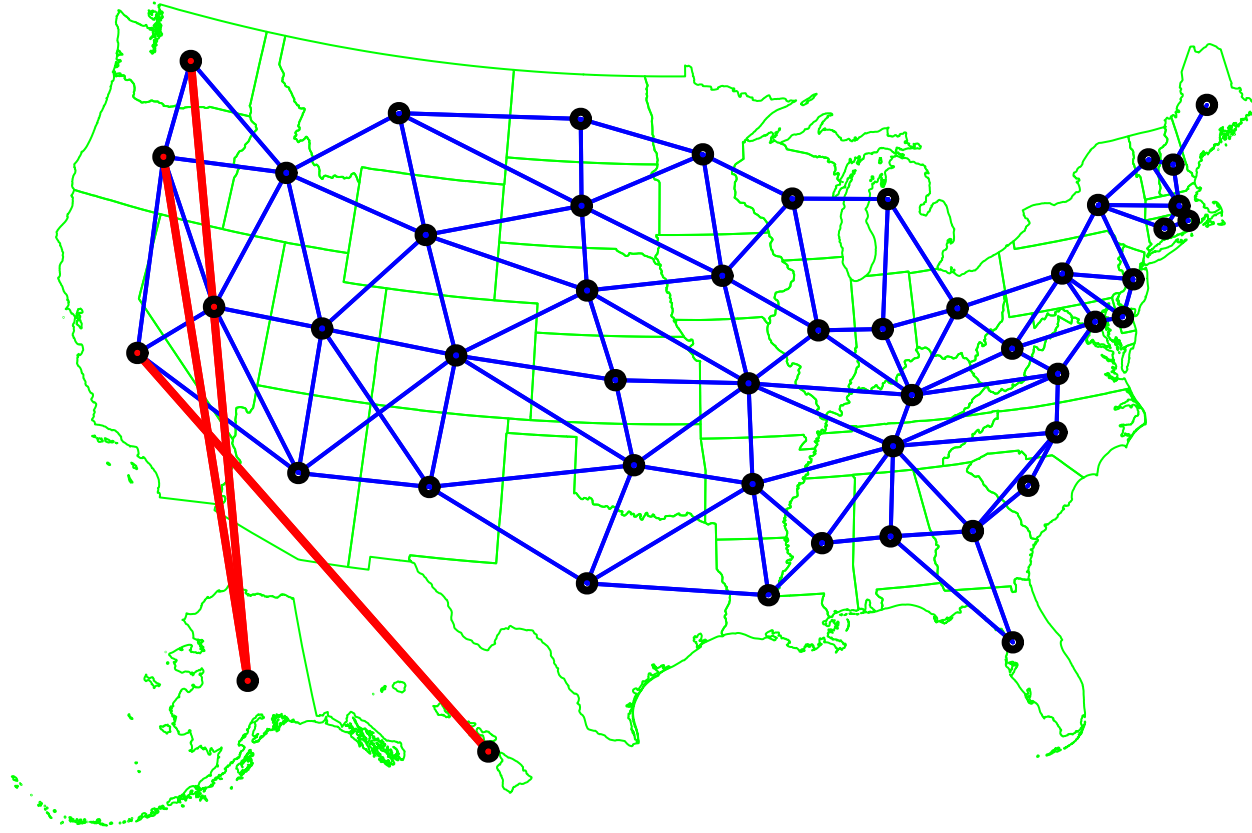
Log
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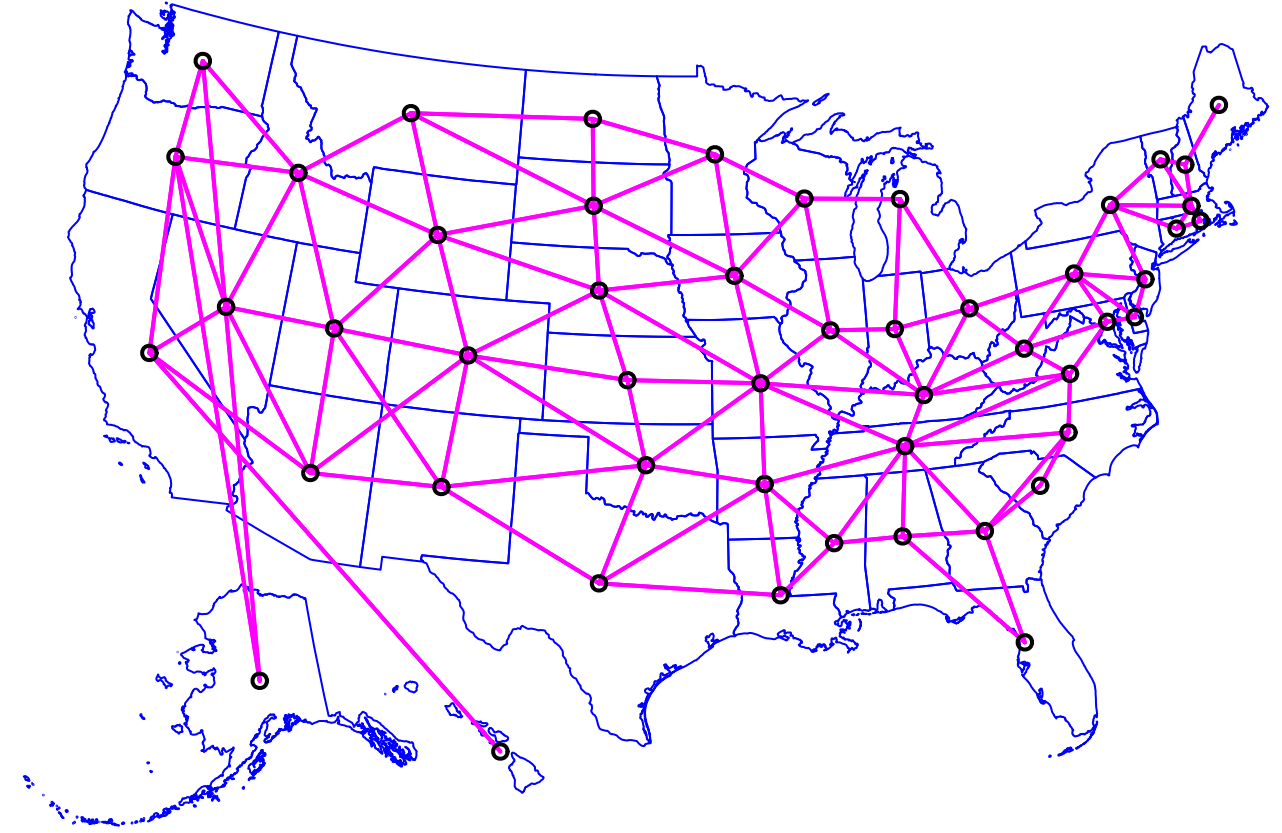
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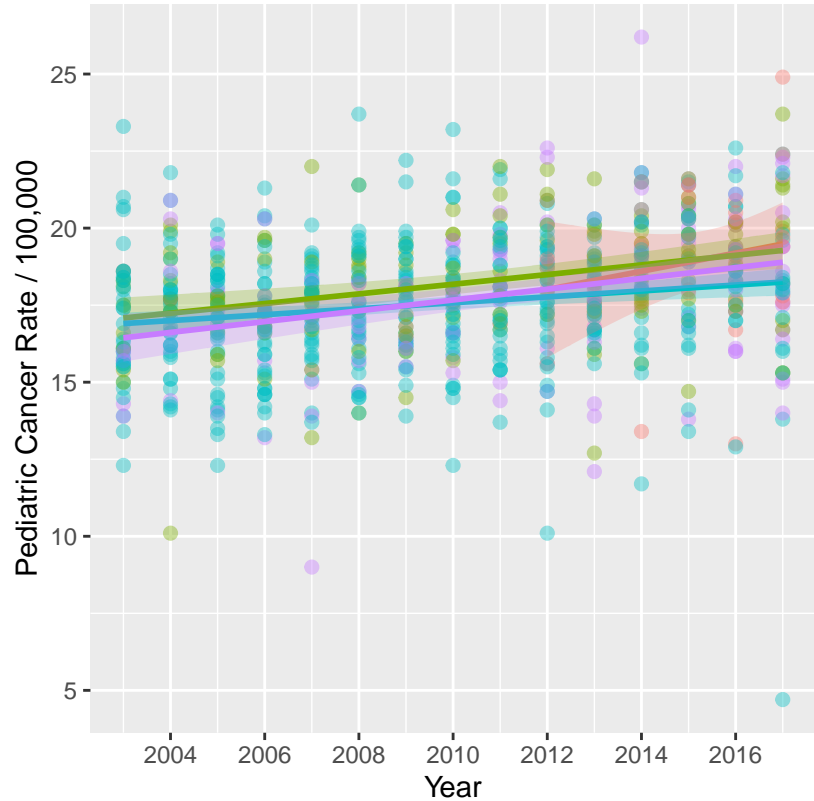
**Additional Links to State Neighbourhood Linkage Network
for US Pediatric Cancer Network Dataset**



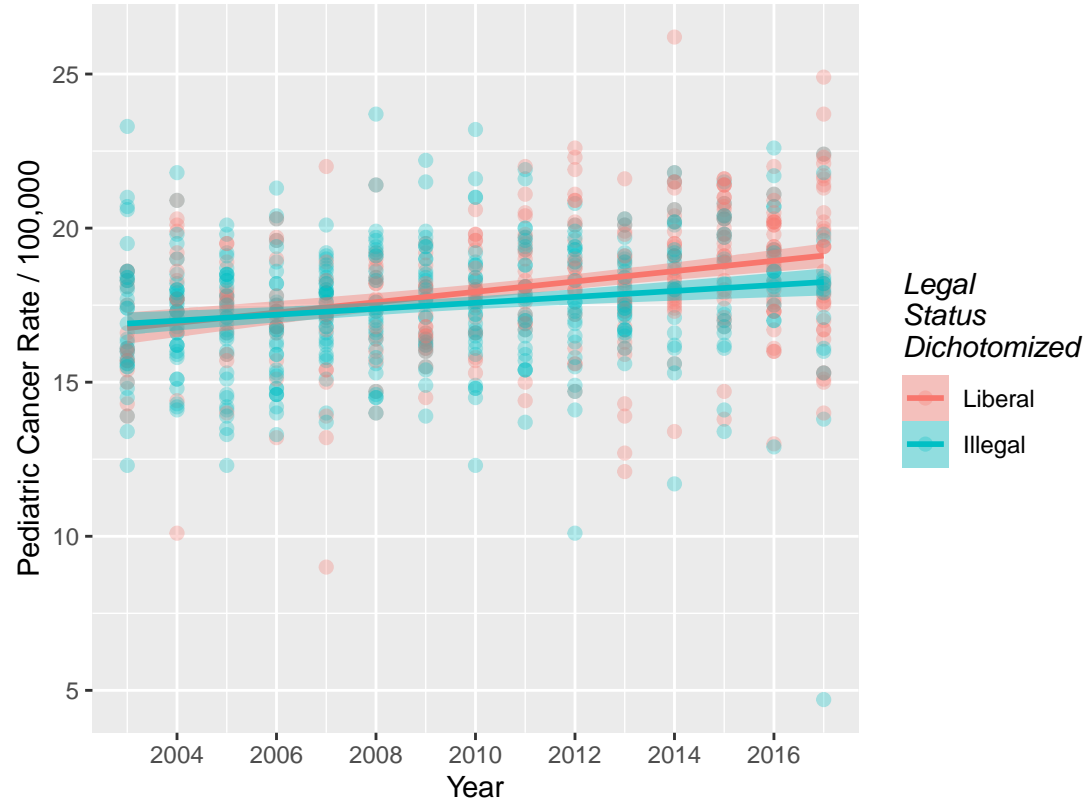
**Final Links to State Neighbourhood Linkage Network
for US Pediatric Cancer Network Dataset**



Cancer Rates by Cannabis Use Legal Status All Statuses



Cancer Rates by Cannabis Use Legal Status Dichotomized – Illegal v Others



Supplementary Table 1.: Cannabis Quintile Data

Quintile	Cannabis Exposure	Cancer Rates
Quintiles		
Quintile 1	0.1101 (0.0038)	17.2941 (0.1913)
Quintile 2	0.1349 (0.0042)	17.4933 (0.1763)
Quintile 3	0.1552 (0.0044)	17.4381 (0.1806)
Quintile 4	0.1731 (0.005)	18.0087 (0.1581)
Quintile 5	0.2304 (0.0062)	18.6060 (0.1767)
Dichotomized Quintiles		
Lower Quintiles	0.1331 (0.0025)	17.4076 (0.1055)
Upper Quintiles	0.2018 (0.0043)	18.3073 (0.1196)

Selected and Major Paediatric Cancer Rates Over Time 21 US Cancer Registries

