

# Context-dependent selection as the keystone in somatic evolution of cancer

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## Abstract

Somatic evolution of cancer involves a series of mutations, and attendant changes, in one or more clones of cells. Unlike a “bad luck” type model, the notion of clonal expansion adds competition-driven selection to the supposedly random process of somatic mutagenesis, with the implicit assumption that any mutation leading to partial loss of regulation of cell proliferation will give a selective advantage to the mutant. However, a number of experiments show that an intermediate pre-cancer mutant has only a conditional selective advantage; given that tissue microenvironmental conditions differ across individual organisms, this selective advantage to a mutant should be widely distributed over the population of organisms. We evaluate three models, namely “bad luck”, context-independent, and -dependent selection, in a comparative framework, on their ability to predict patterns in total incidence, age-specific incidence, and their ability to explain Peto’s paradox. Results show that context dependence is necessary and sufficient to explain observed epidemiological patterns, and that cancer incidence is largely selection-limited, as opposed to the mutation-centric, “bad luck” view. A wide range of physiological, genetic and behavioural factors influence the tissue micro-environment, and could therefore be the source of this context dependence in somatic evolution of cancer. The identification and targeting of these micro-environmental factors that influence the dynamics of selection offer new possibilities for cancer prevention. Our work also seeks to renew interest in the comparative evaluation framework, whose application has seen a lull in cancer literature, despite the possibilities of rejection it offers for potential theories of carcinogenesis.

## 1 Epidemiological observations

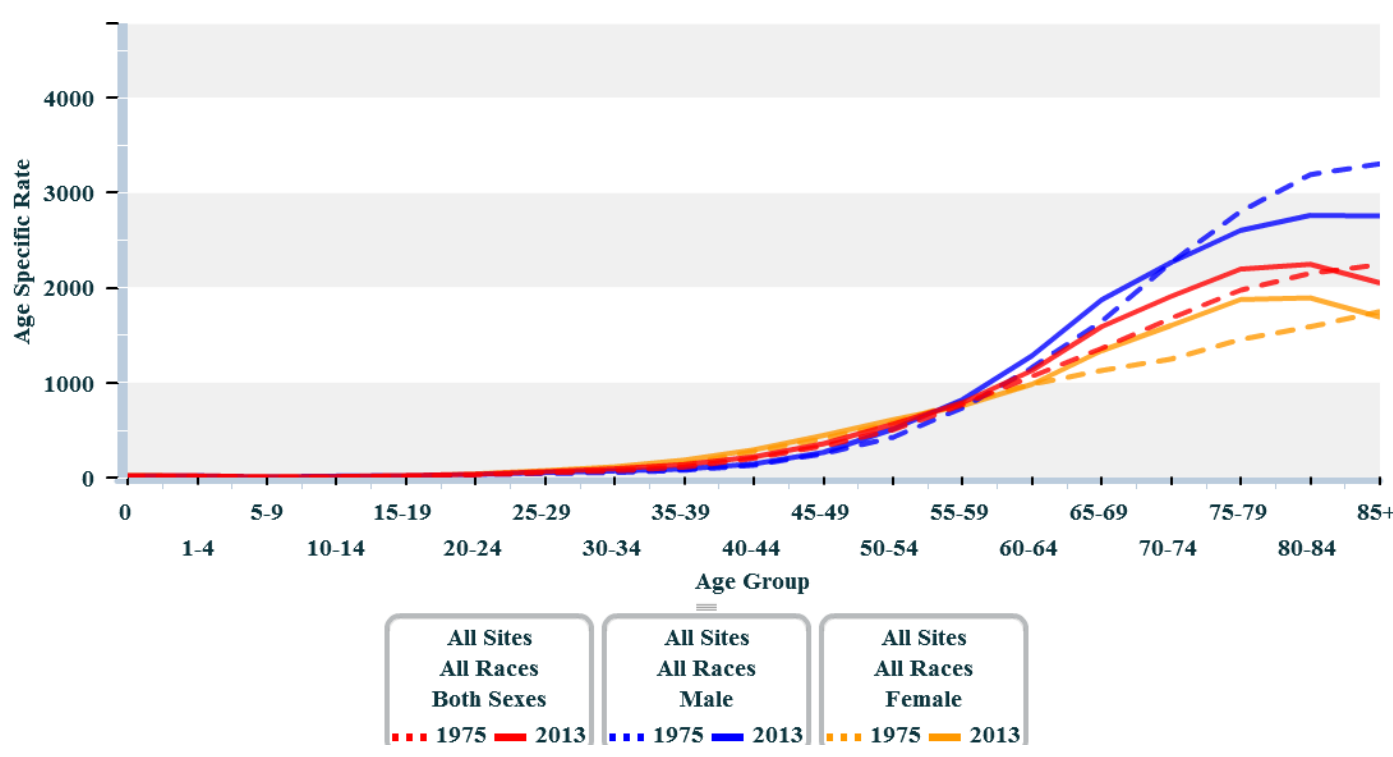


Figure 1: Cancer incidence vs age, from SEER9 (1)

## 2 “Bad luck” model

Equations used:

- $p_{can} = 1 - (1 - p)^n$
- $p_A = 1 - (1 - p_{can})^A$

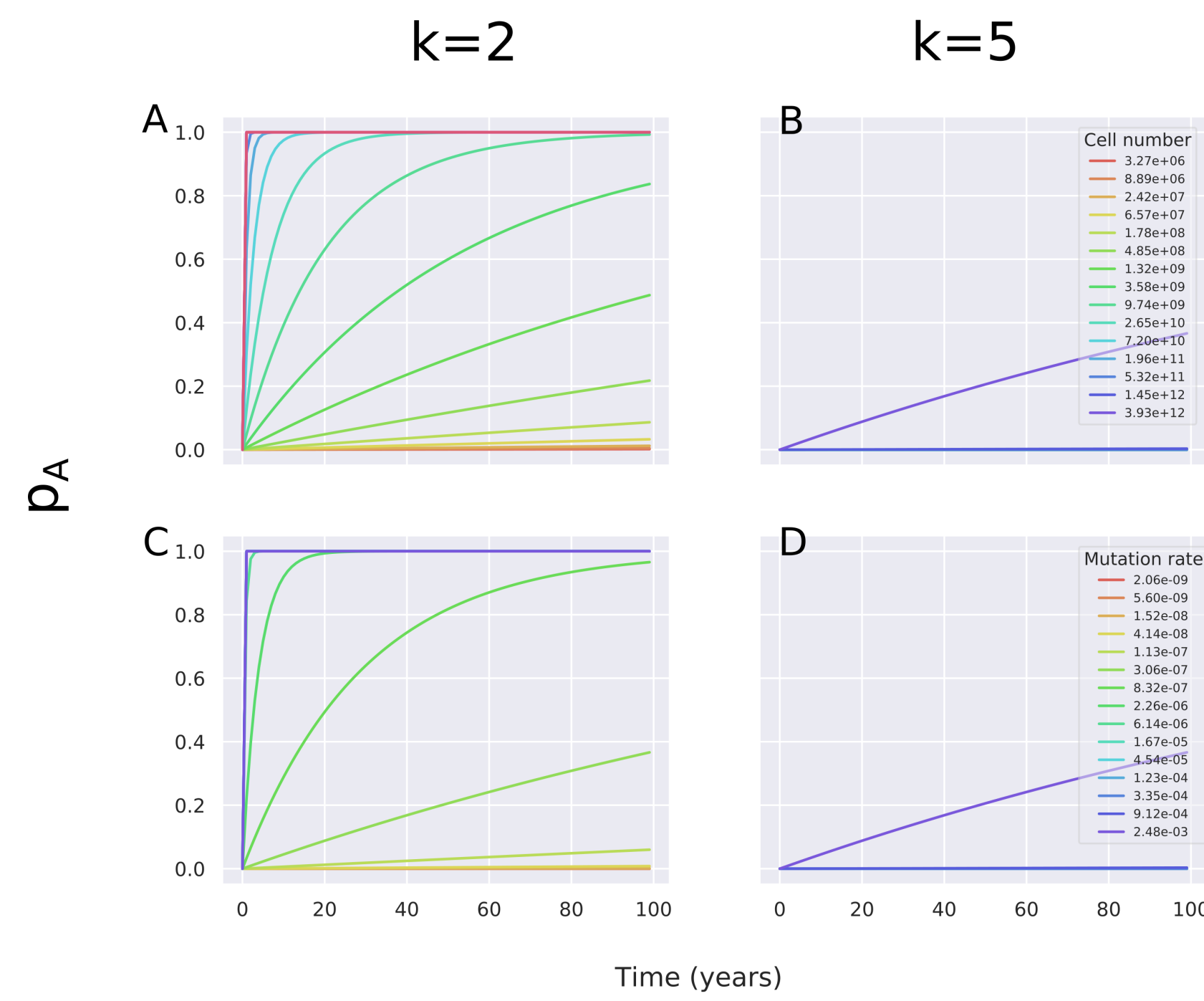


Figure 2: Incidence vs age, with  $n$  and  $p$

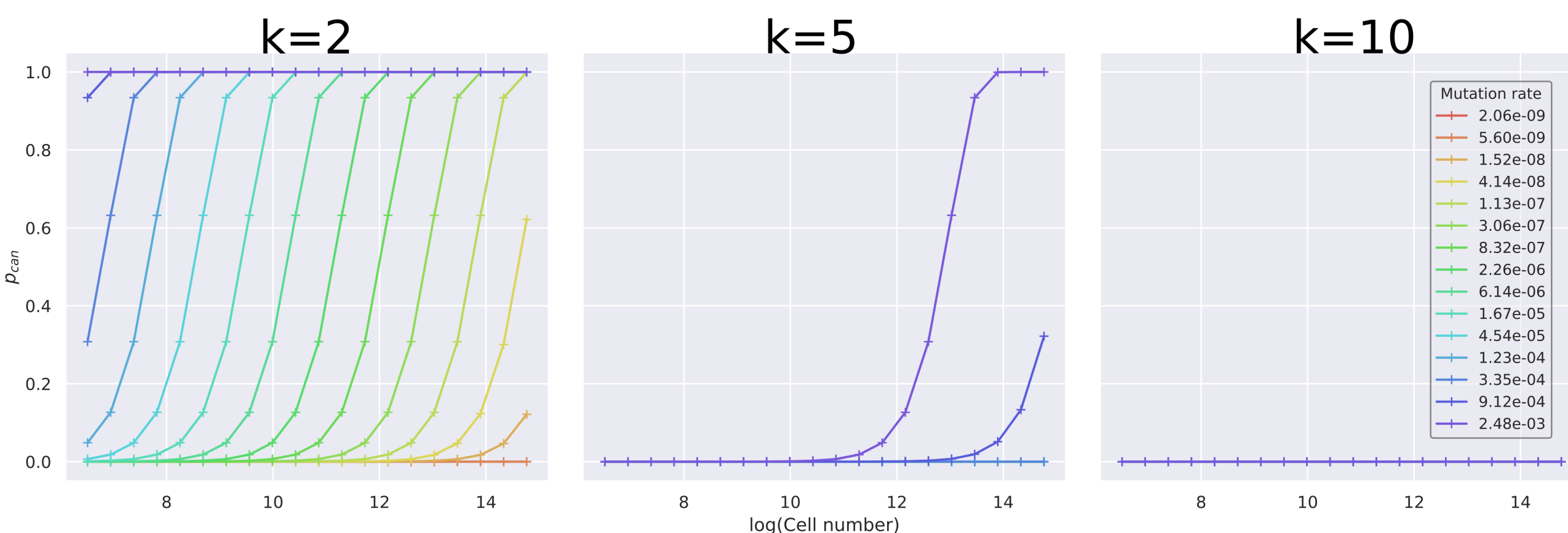


Figure 3: Incidence vs  $\log(n)$  for different  $p$  and  $k$

## 3 Selection models

- Linear evolution process, leading to mutation accumulation
- Discrete logistic equation for cell growth and competition, with one step growth making one day of lifespan
- Non-mutant growth rate,  $g_0 = 0.007$ , and a linear progression up to  $g_k$  for the  $k$ th oncogenic mutation, at which cancer occurs
- $\Delta g = \frac{g_k - g_0}{k}$  is randomized in the population for context-dependent selection, as  $N(\mu, \sigma)$
- $n \in [1.203 * 10^6, 2.649 * 10^{10}]$
- $p \in [3.775 * 10^{-11}, 3.059 * 10^{-7}]$

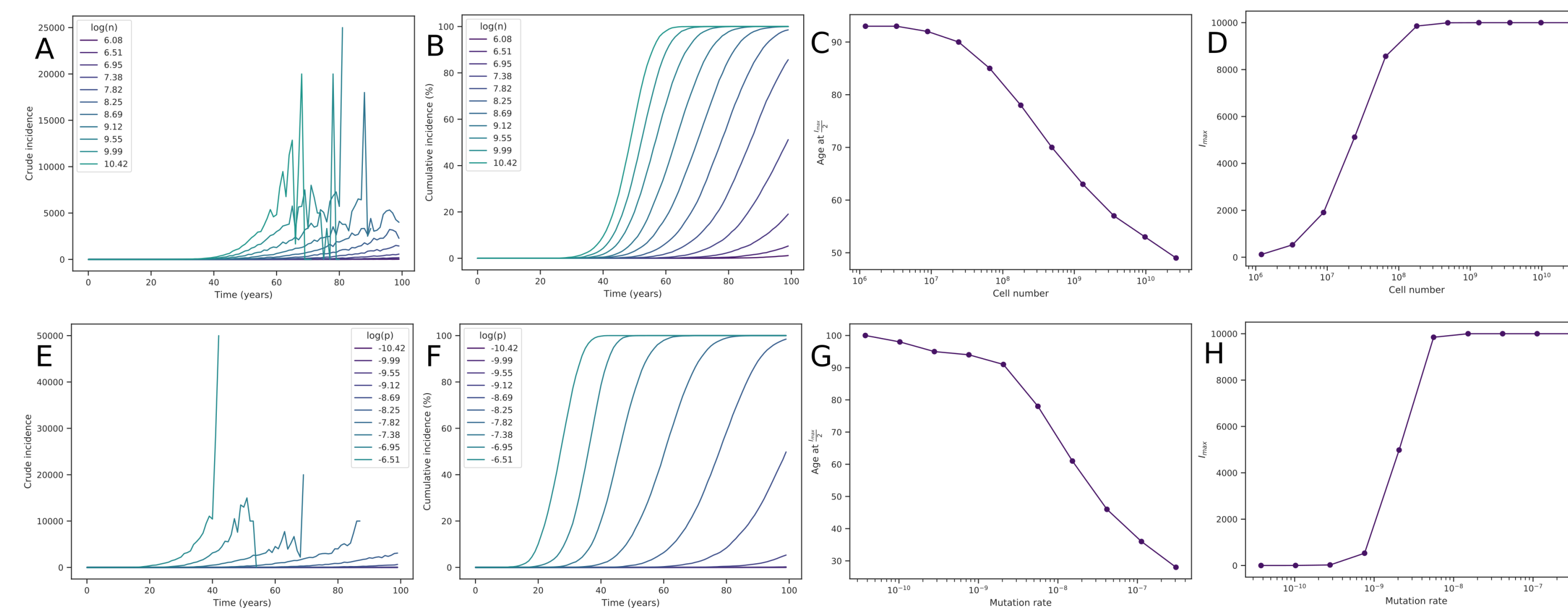


Figure 5: Context-independent selection; crude and cumulative incidence rates vs age, age at half maximum cumulative incidence, and maximum cumulative incidence, over the range of  $n$  and  $p$ .

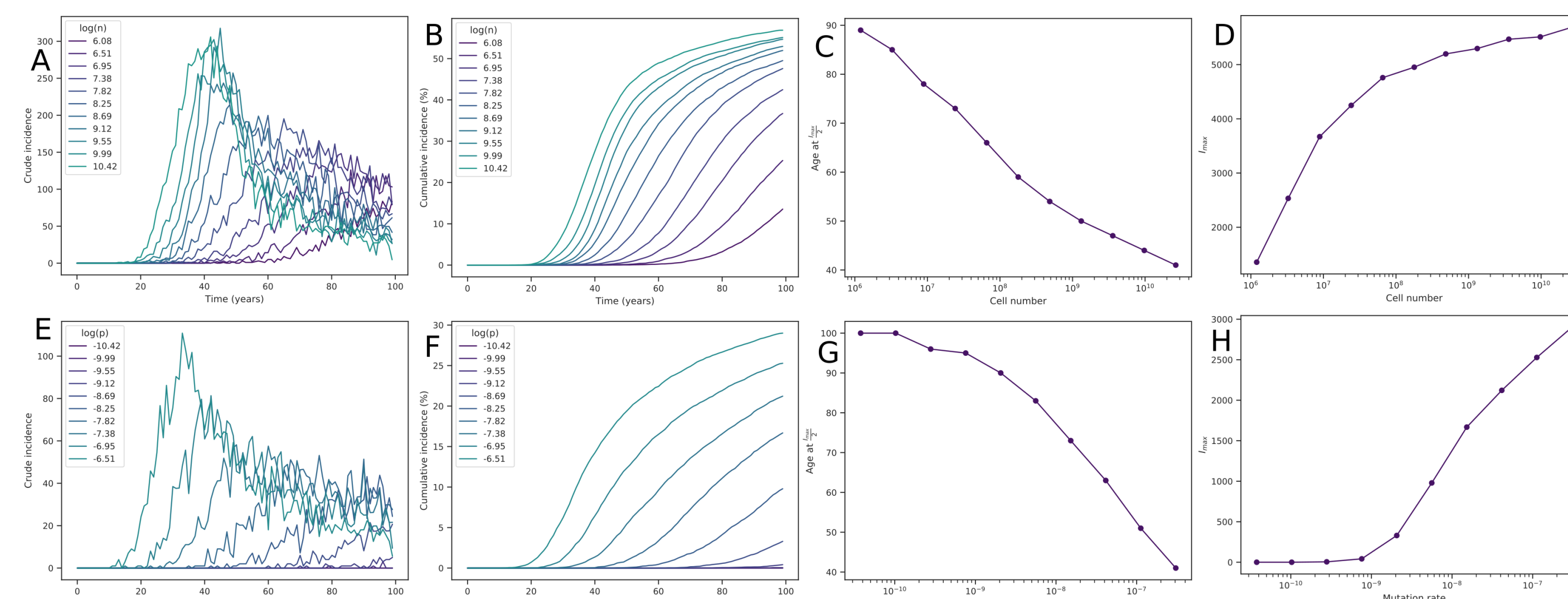


Figure 6: Same incidence parameters as above, for the context-dependent selection case.

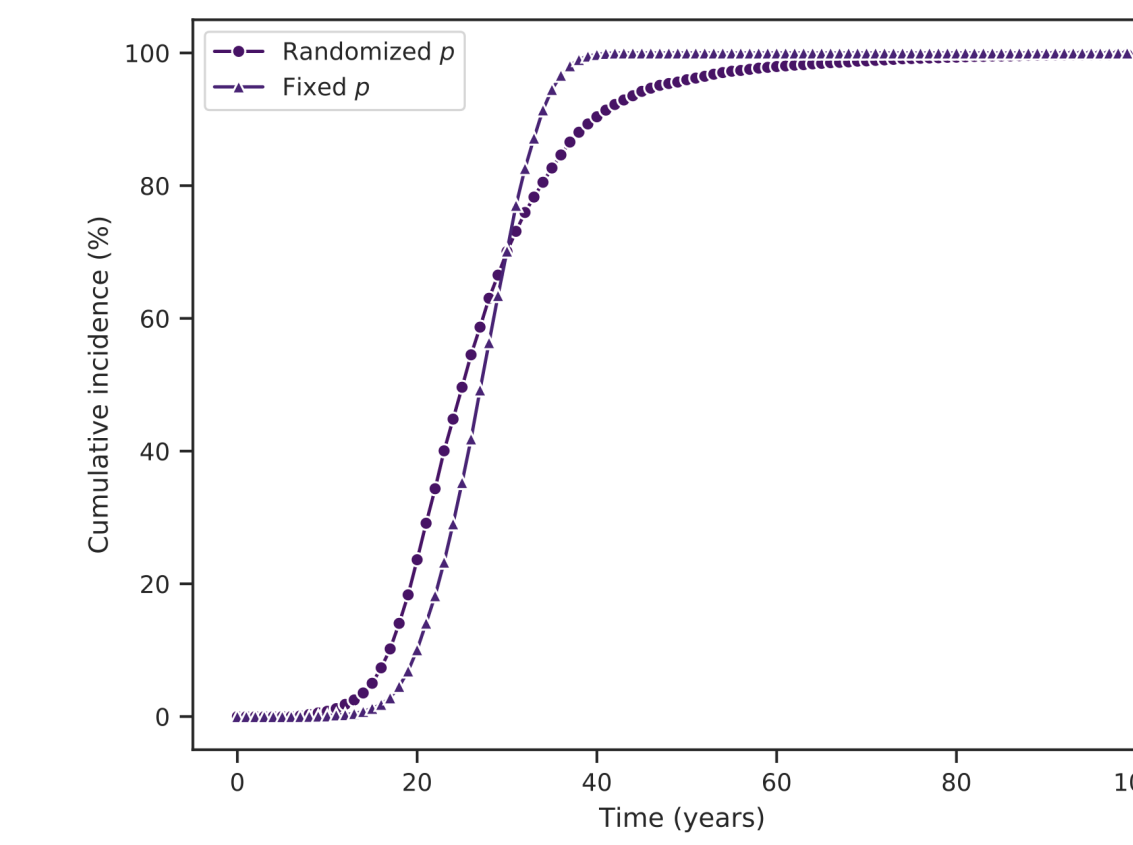


Figure 4: Context-independent model; randomized  $p$  vs fixed  $p$ . Mean randomized  $p \approx$  fixed  $p$ .

### 3.1 Effect of $k$

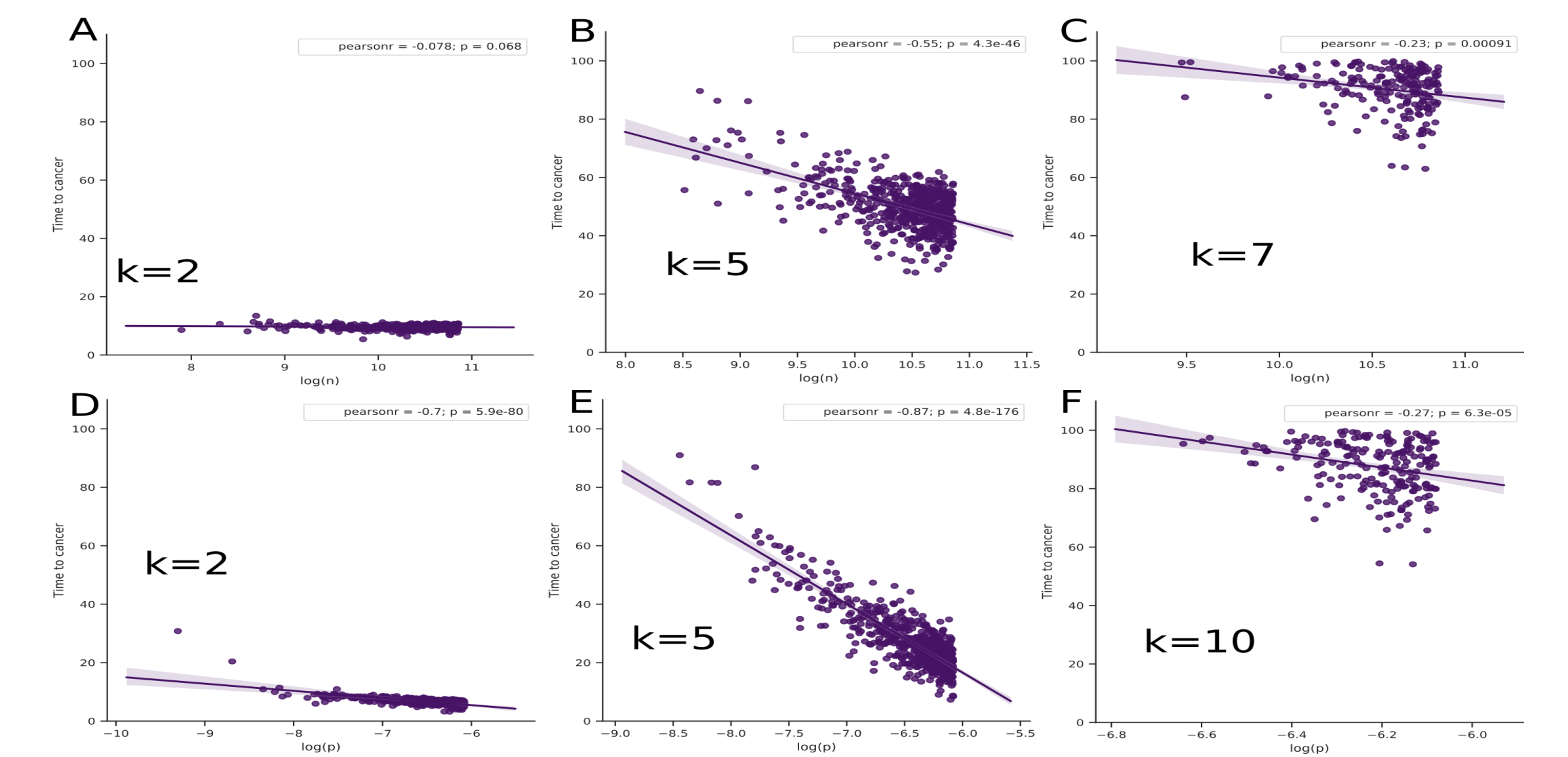


Figure 7: Context-independent selection case; association of time to cancer with  $n$  and  $p$  is modulated by  $k$ .

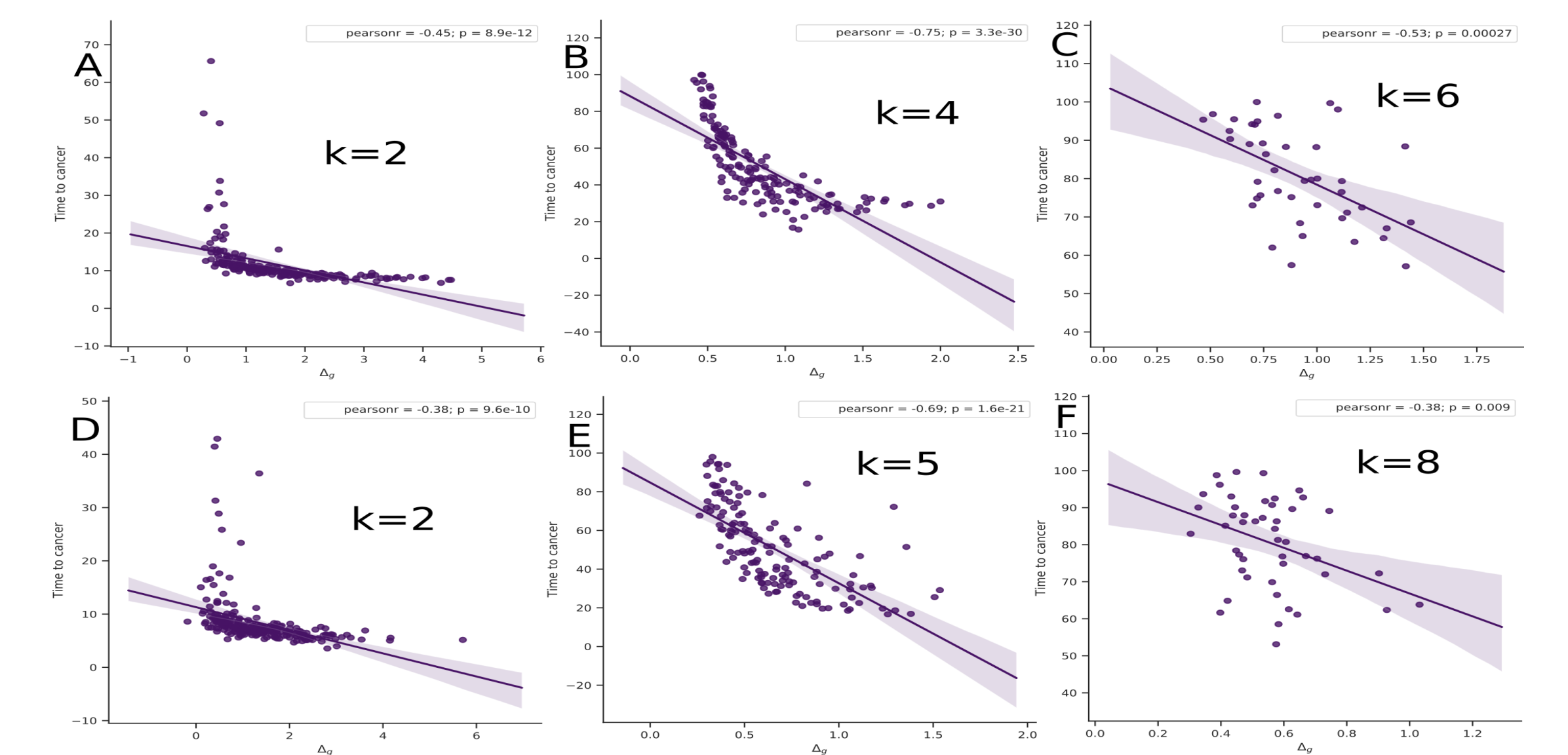


Figure 8: Context-dependent selection case; association of time to cancer with  $\Delta g$  is modulated by  $k$ .

## 4 Conclusions

Epidemiological observation	“Bad luck”	Context-independent selection	Context-dependent selection
Total incidence < 30%	100%	100%	<100% possible
Late-life decline	No	No	Yes
Incidence vs $n$	Threshold	Threshold	Progressive
Non-mutagenic carcinogens	Incompatible	Incompatible	$g$ distribution
Peto’s paradox	Extrinsic <sup>1</sup>	Extrinsic <sup>1</sup>	Intrinsic

## References

- (1) American Cancer Society. “Cancer Facts & Figures 2016”. In: Cancer Facts & Figures 2016 (2016), pp. 1–9.

## Acknowledgements

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