

1	Doses under investigation?	
2	Target toxicity level (TTL)?	
3	<p>Skeleton?</p> <p>If no skeleton, which dose do you expect to be the MTD? You can use <code>dfcrm::getprior</code> to generate a prior. E.g.  <code>getprior(halfwidth = 0.05, target = 0.2, nu = 4, nlevel = 5)</code>  will generate prior that anticipates dose-level 4 of 5 is the sought dose with associated <math>\text{Prob}(\text{DLT}) = 0.2</math>.  Tweak <code>halfwidth</code> to get a prior that you agree with.</p>	
4	<p>Starting dose?</p> <p>This might be lower than your guess at MTD.  Do you have doses to de-escalate to, if your assumptions are wrong? Painful to stop a trial due to poor planning.</p>	
5	Model type?	
6	Model parameters, including prior hyperparameters?	
7	<p>How to select dose?</p> <p>Describe constraints, like “no skipping in escalation” or “at least two complete negative DLT evaluations before escalation”</p>	
8	<p>How to know when to stop?</p> <p>Describe constraints like “use no more than 30 patients” or “stop early if lowest dose is too toxic”</p>	
9	Length of DLT assessment window	
	<b>If using non-time-to-event method:</b>	
10	How to select cohort size?	
	<b>If time-to-event method:</b>	
11	How to calculate weight of observation from length of follow-up?	
	<b>For simulation:</b>	

12	What is assumed true $\text{Prob}(\text{Tox})$ ?	
	<b>If time-to-event method:</b>	
13	How to sample time between consecutive patients?	
14	How to sample time of toxicity, given that toxicity happened?	