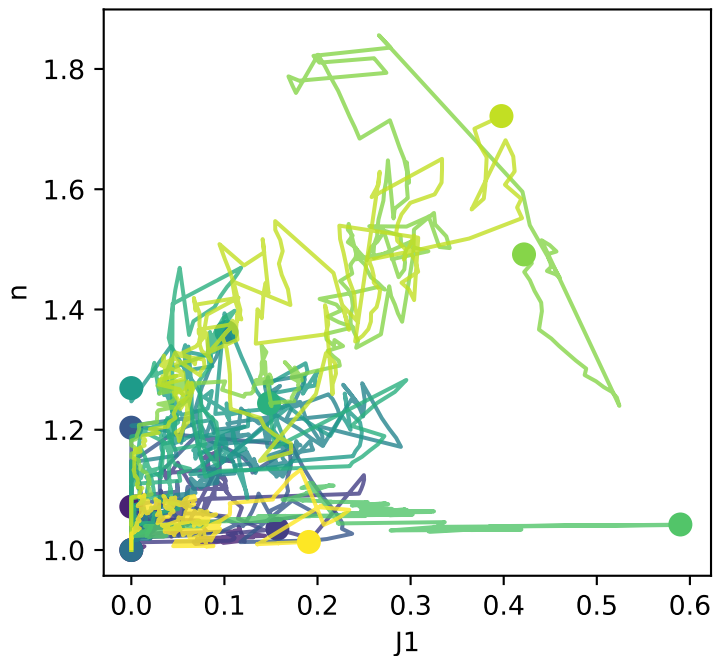
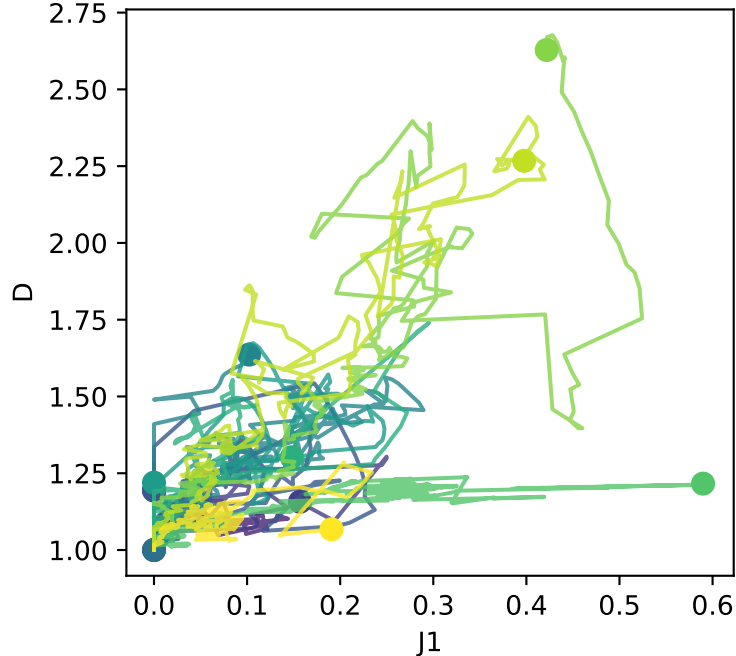


Average trajectories in index space for non-spatial tumours

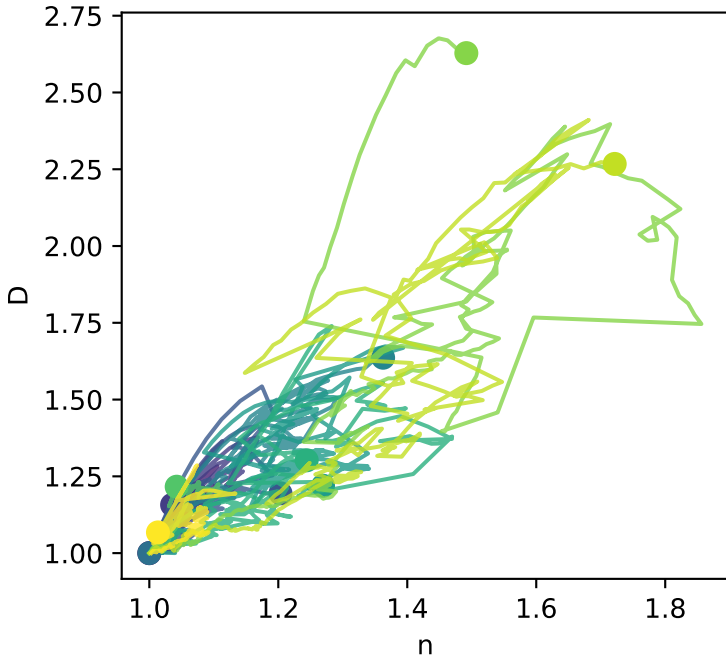
J1 vs n



J1 vs D



n vs D



- Mutation rate, advantage: ('1e-06', 0)
- Mutation rate, advantage: ('1e-06', 0.05)
- Mutation rate, advantage: ('1e-06', 0.1)
- Mutation rate, advantage: ('1e-06', 0.2)
- Mutation rate, advantage: ('1e-05', 0)
- Mutation rate, advantage: ('1e-05', 0.05)
- Mutation rate, advantage: ('1e-05', 0.1)
- Mutation rate, advantage: ('1e-05', 0.2)
- Mutation rate, advantage: ('1e-04', 0)
- Mutation rate, advantage: ('1e-04', 0.05)
- Mutation rate, advantage: ('1e-04', 0.1)
- Mutation rate, advantage: ('1e-04', 0.2)