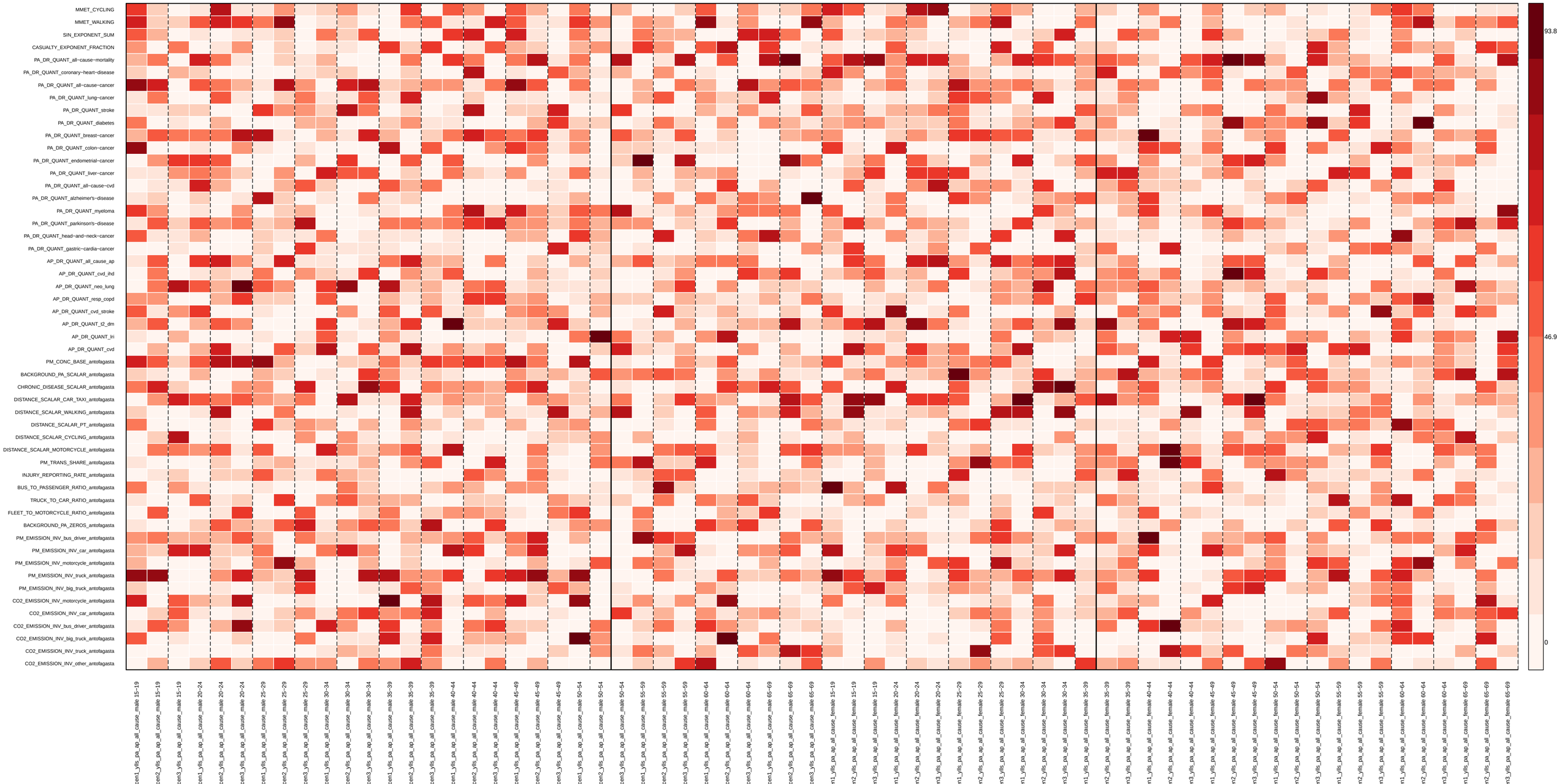
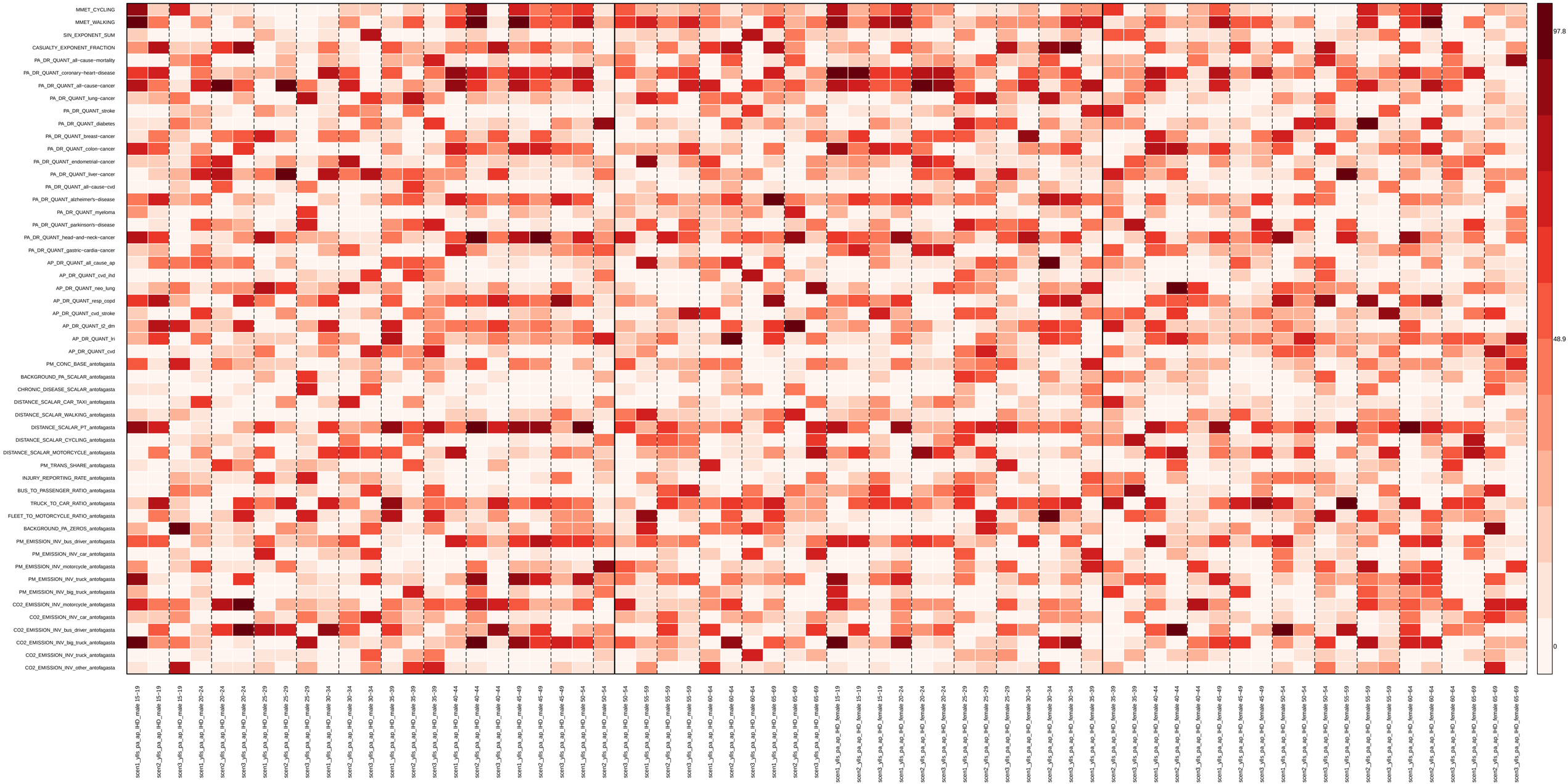
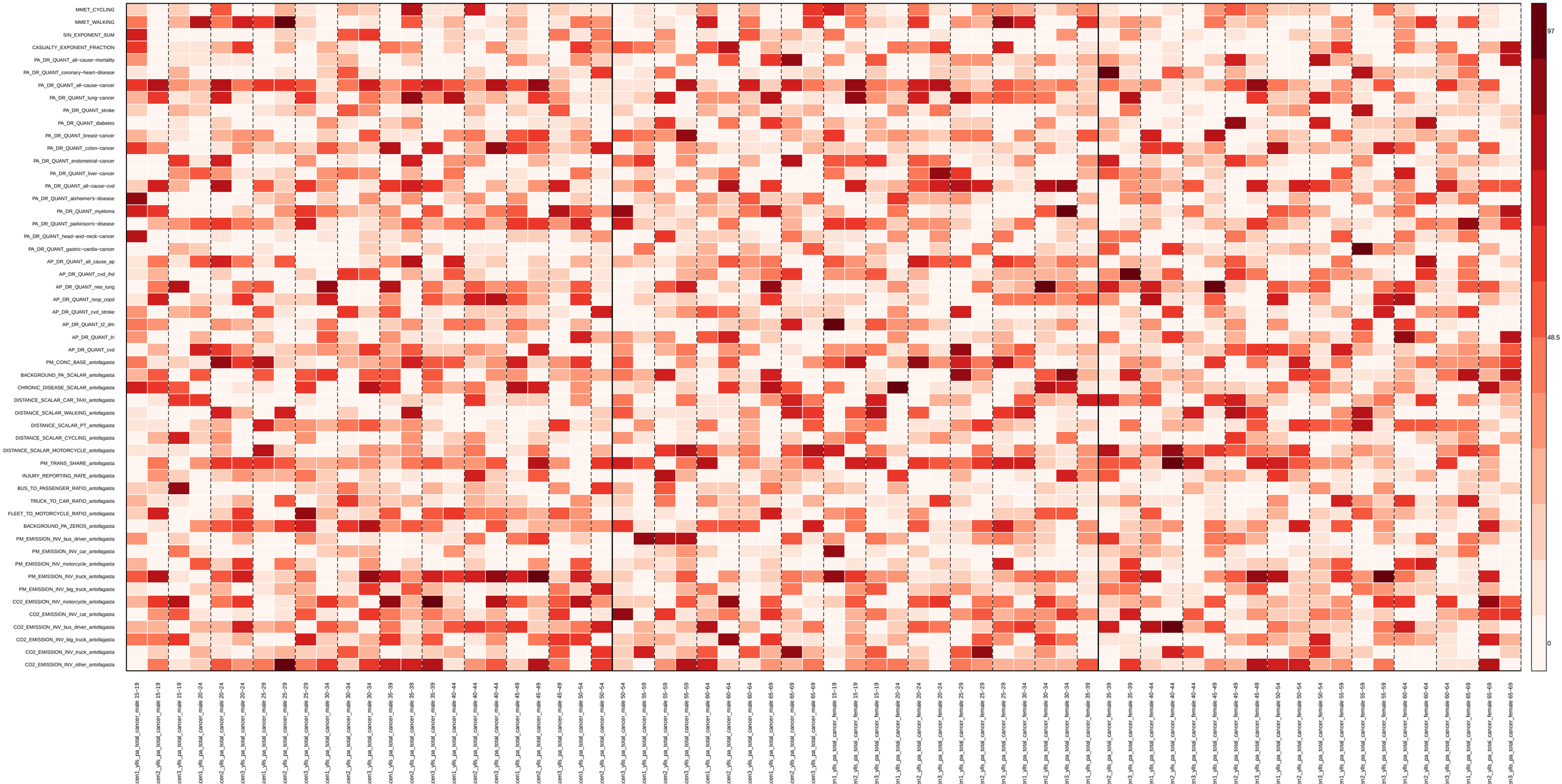
antofagasta − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



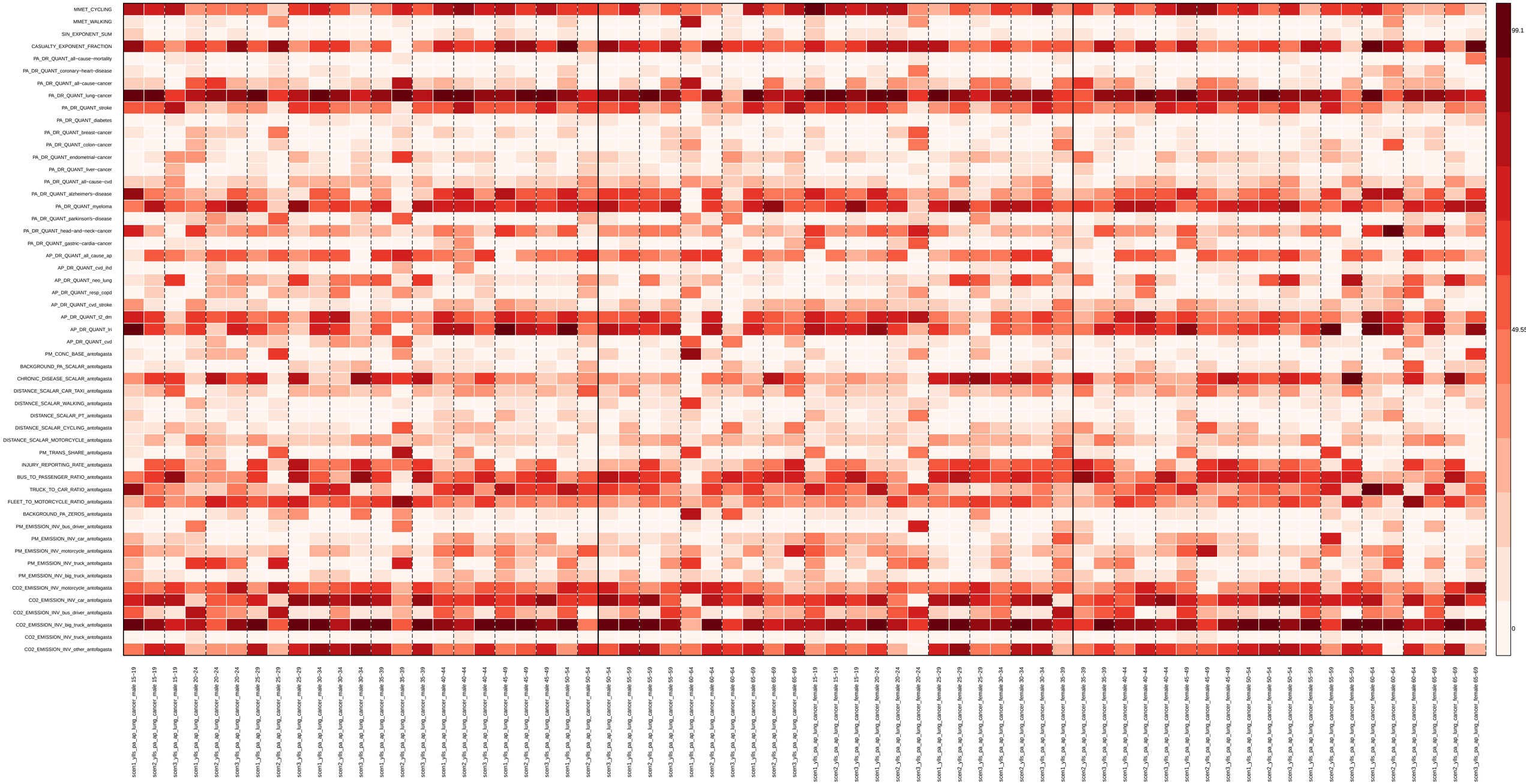
antofagasta − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



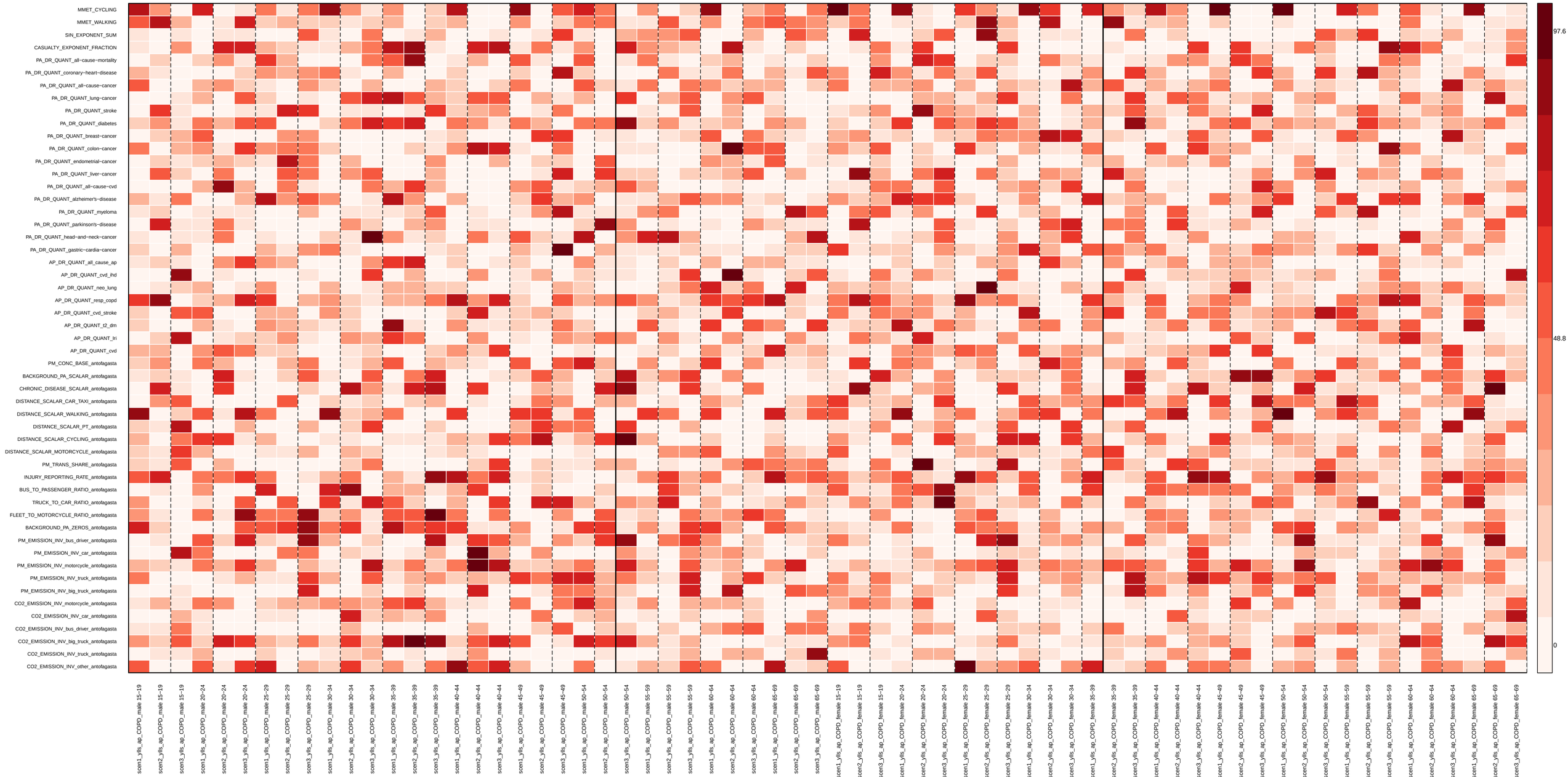
antofagasta − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



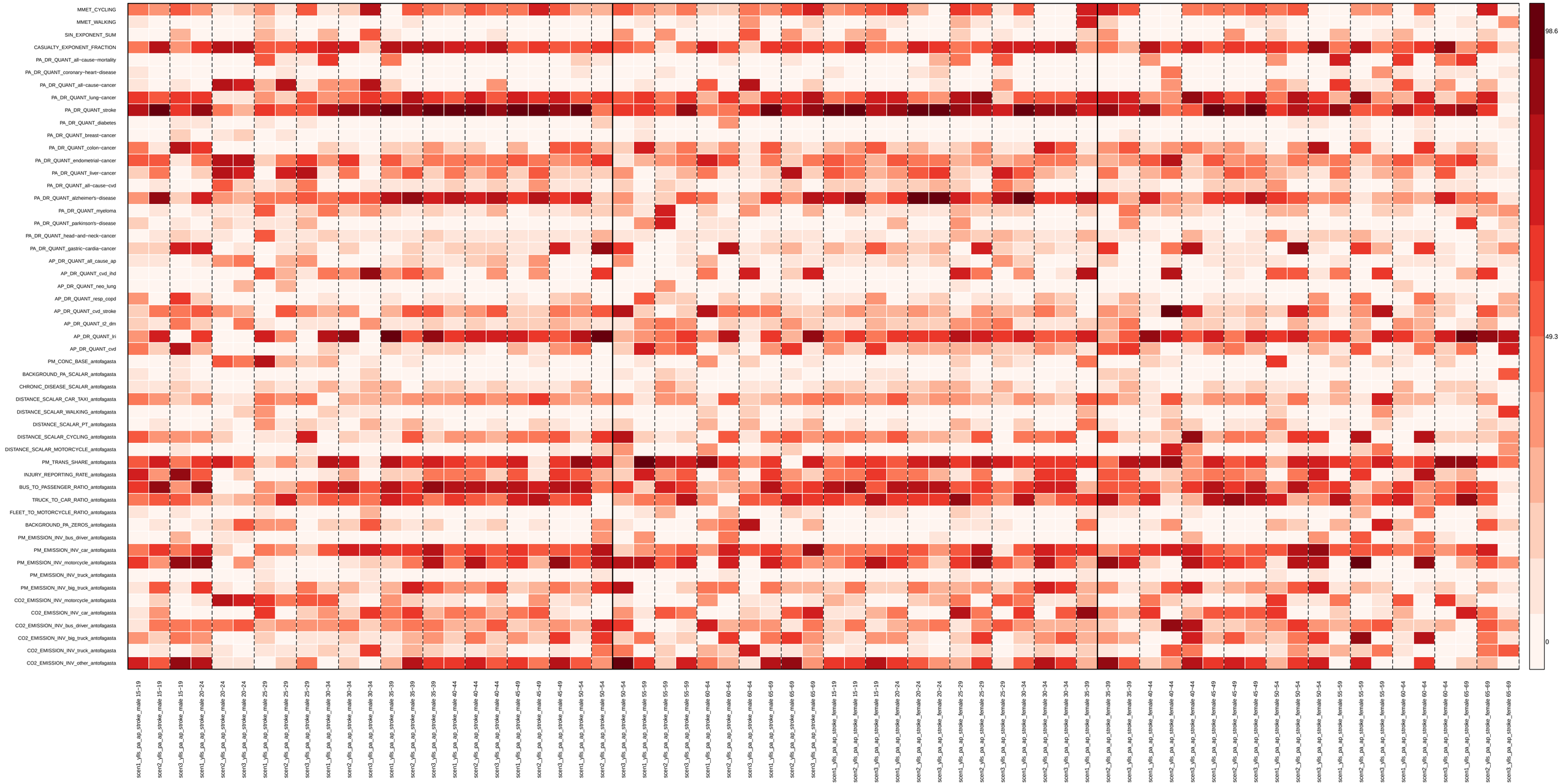
antofagasta − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



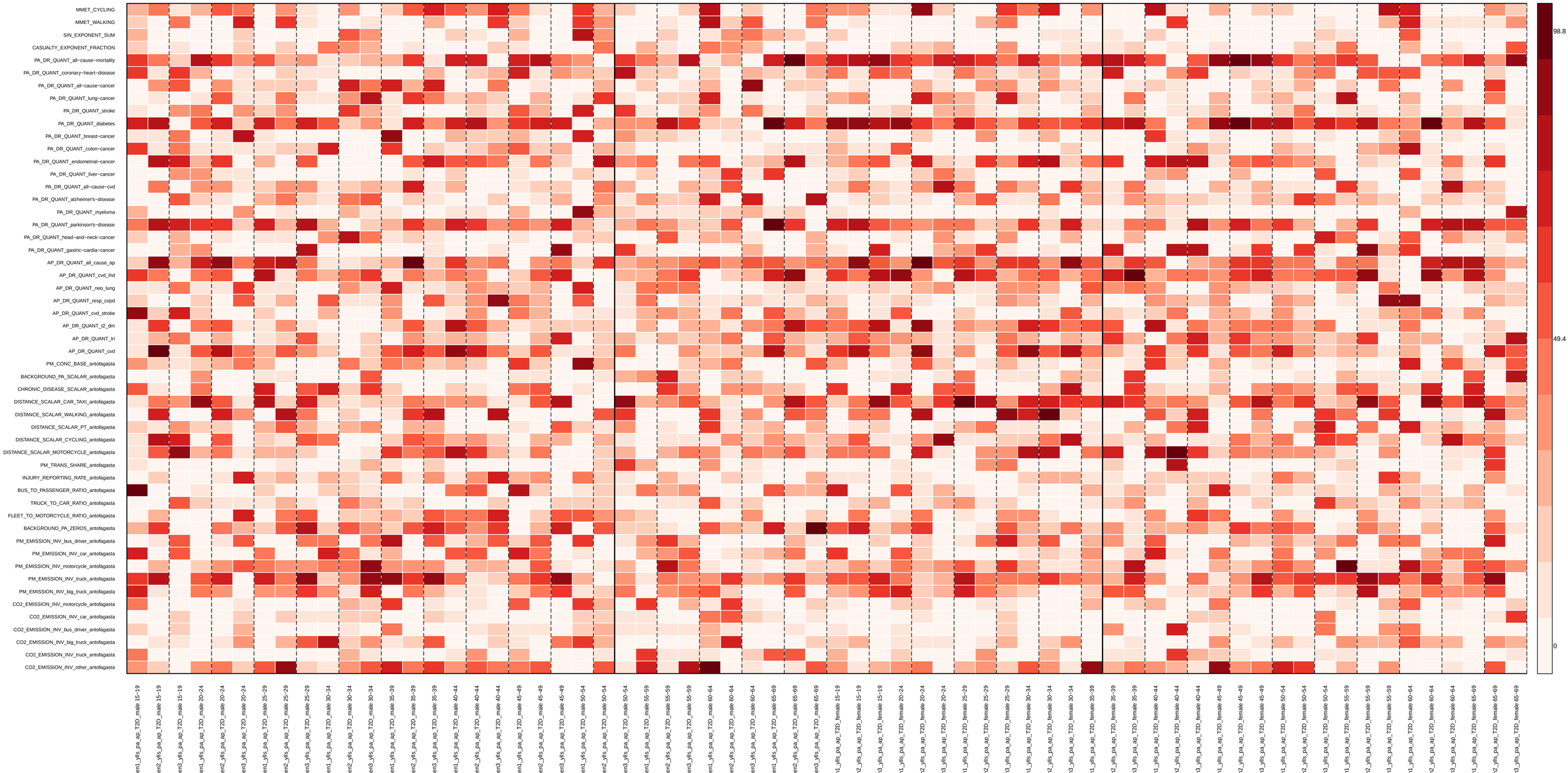
antofagasta − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



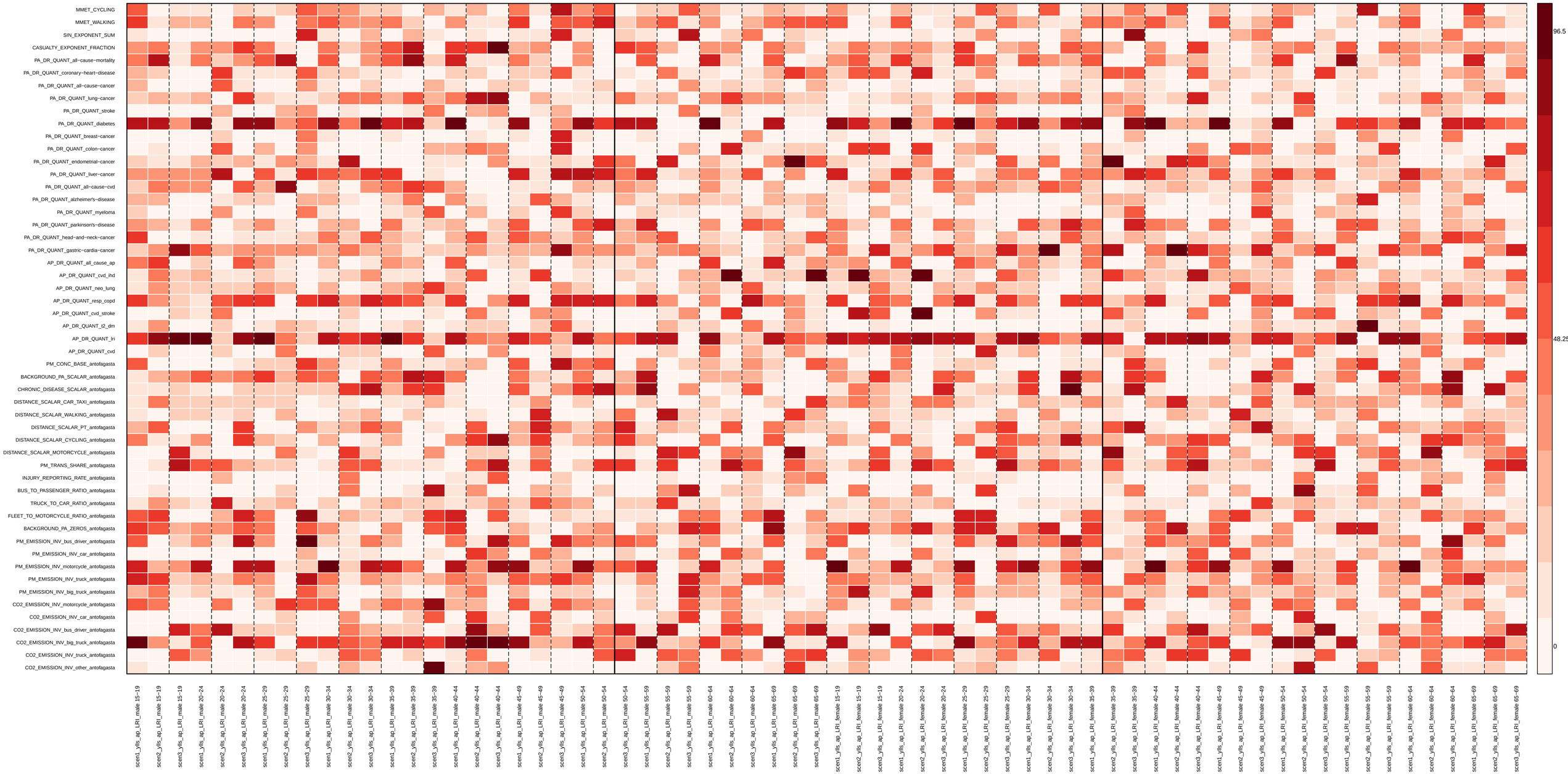
antofagasta − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



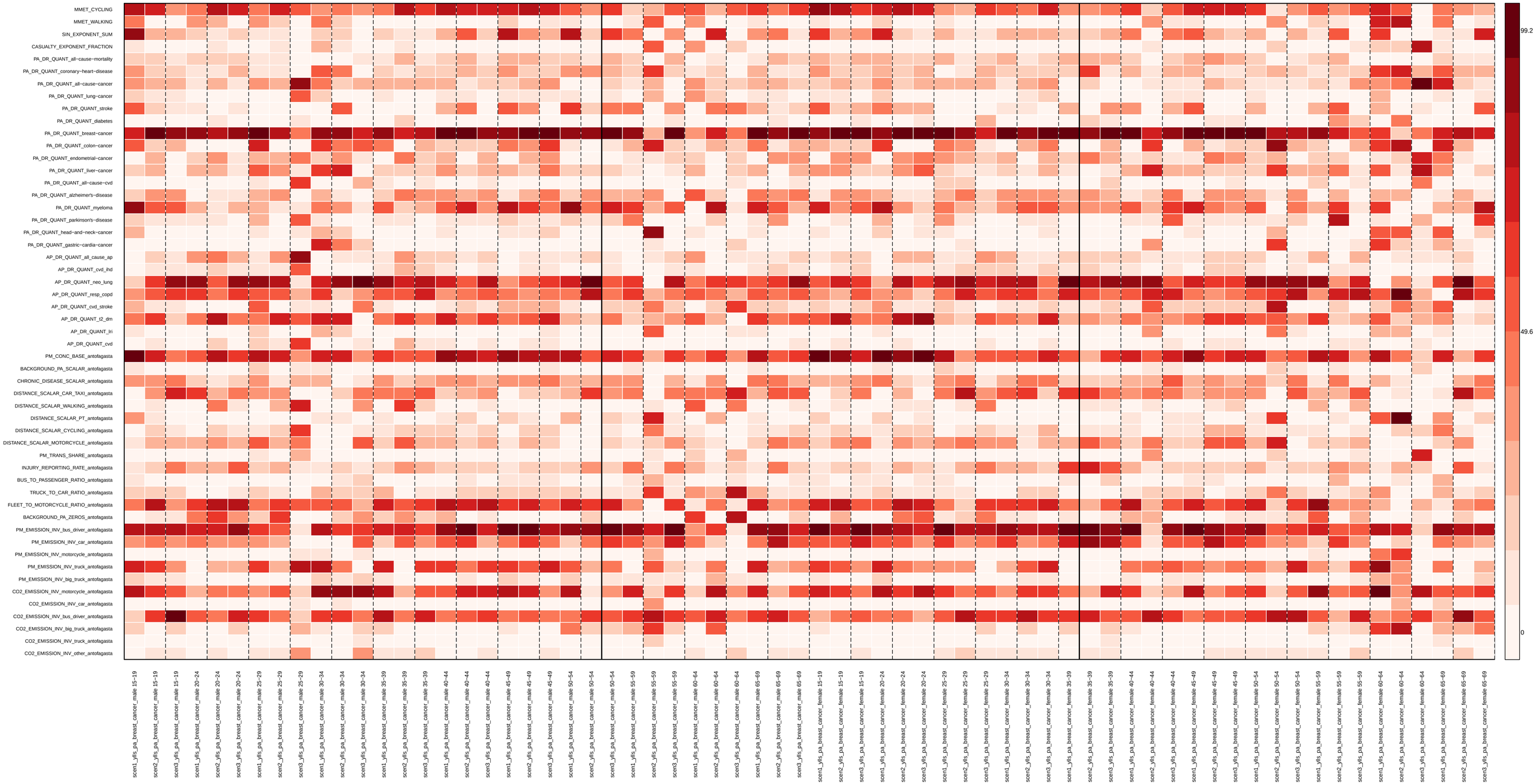
antofagasta − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



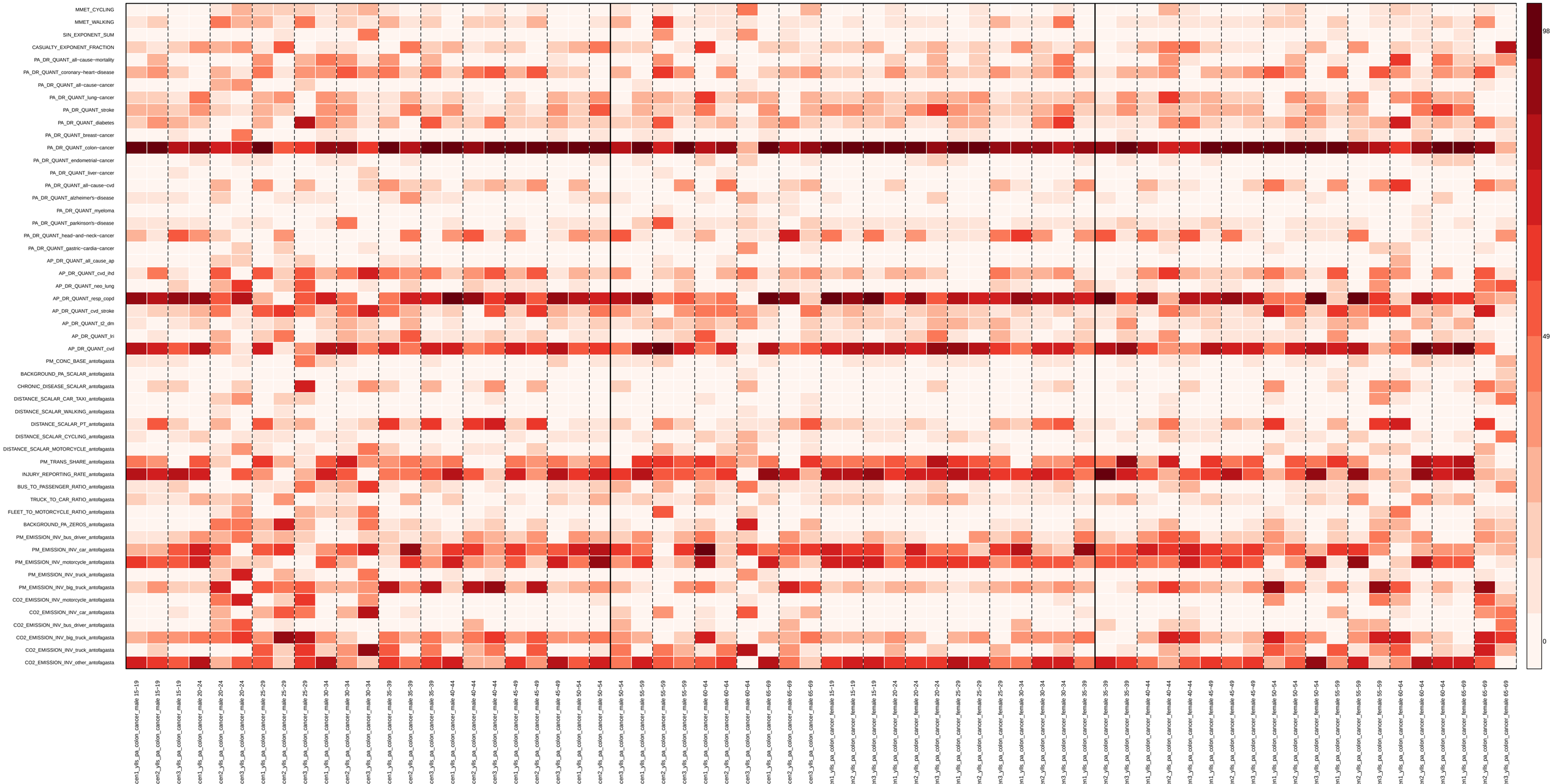
antofagasta − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



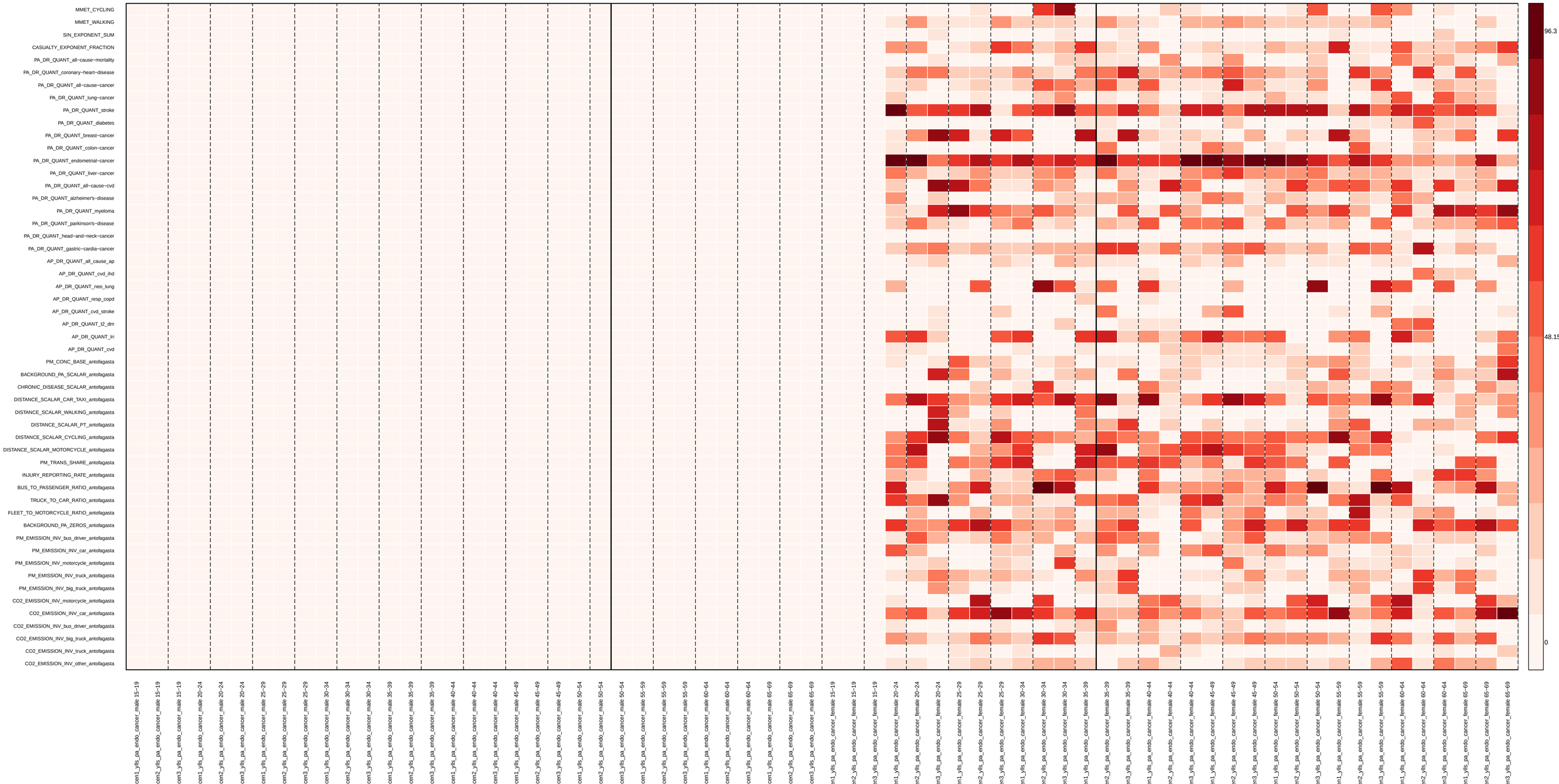
antofagasta − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



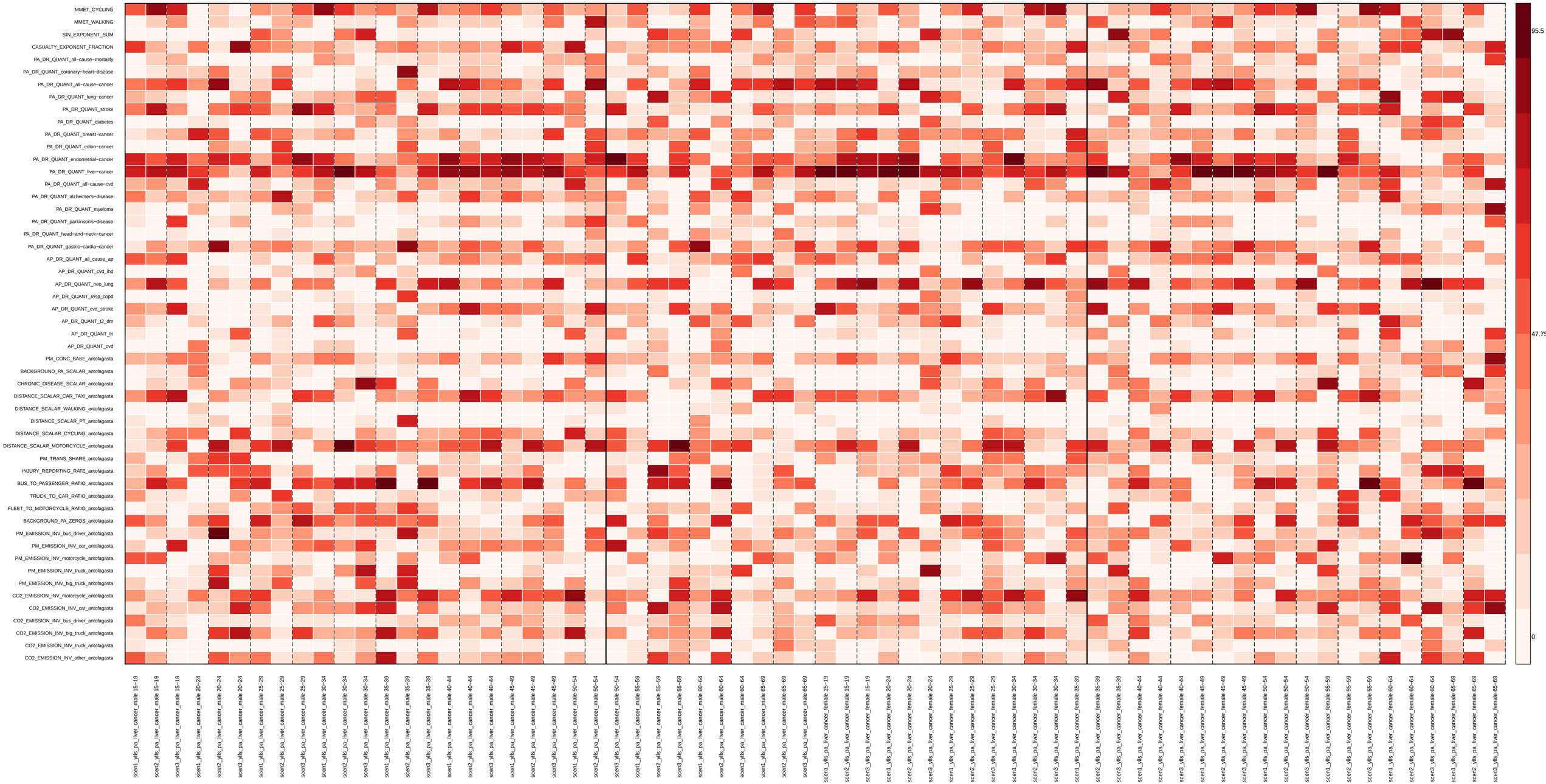
antofagasta − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



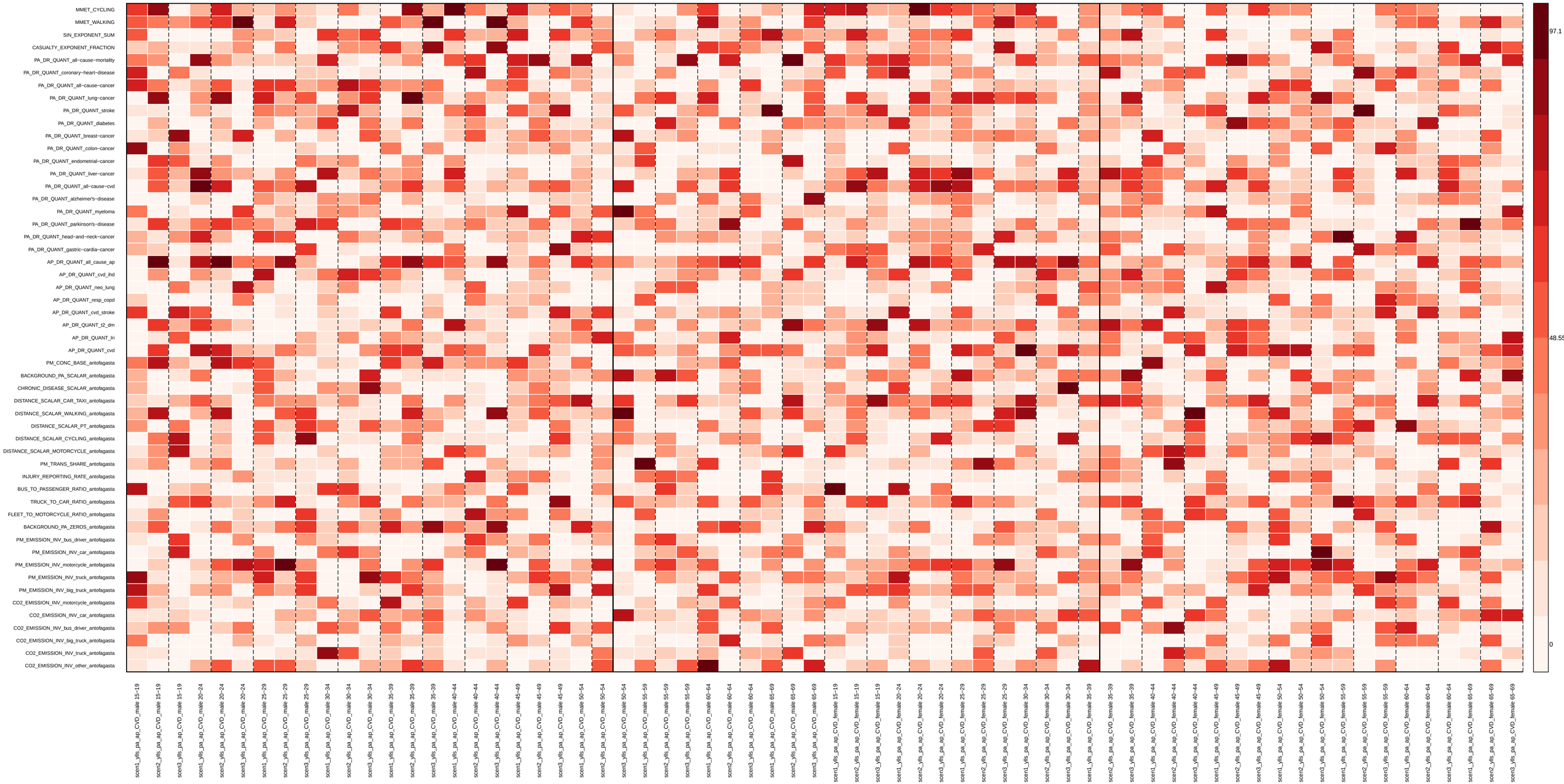
antofagasta − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



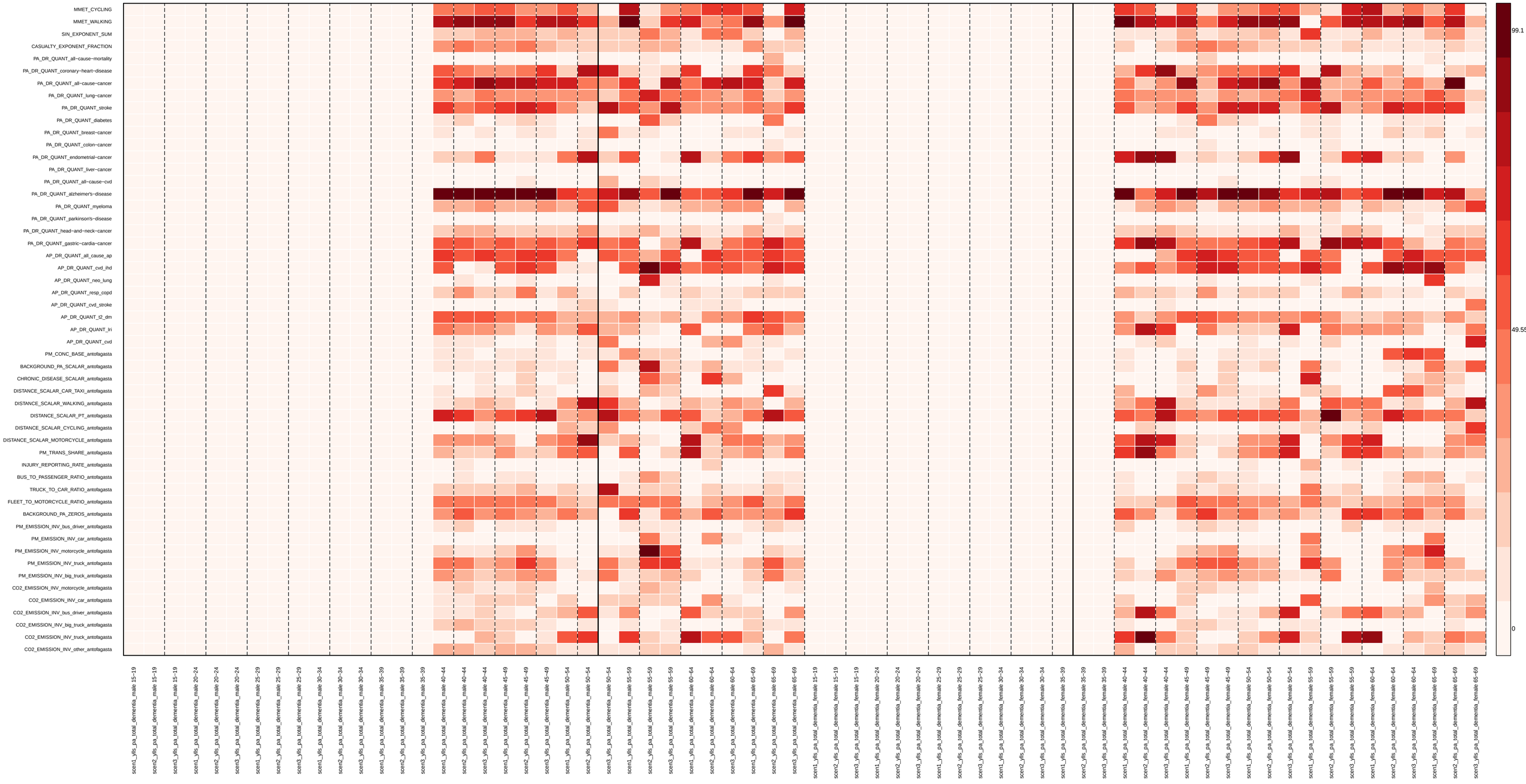
antofagasta − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



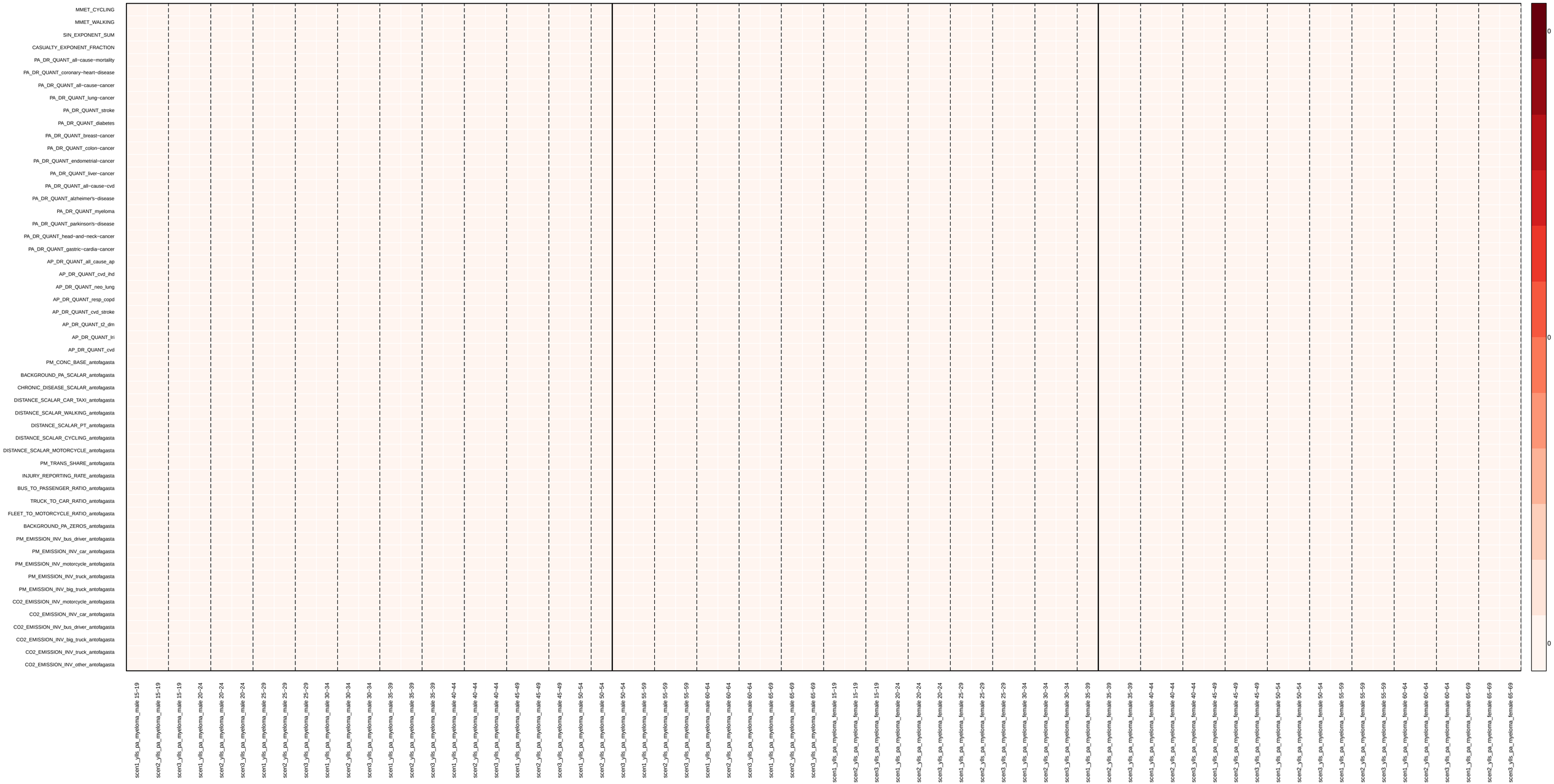
antofagasta − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



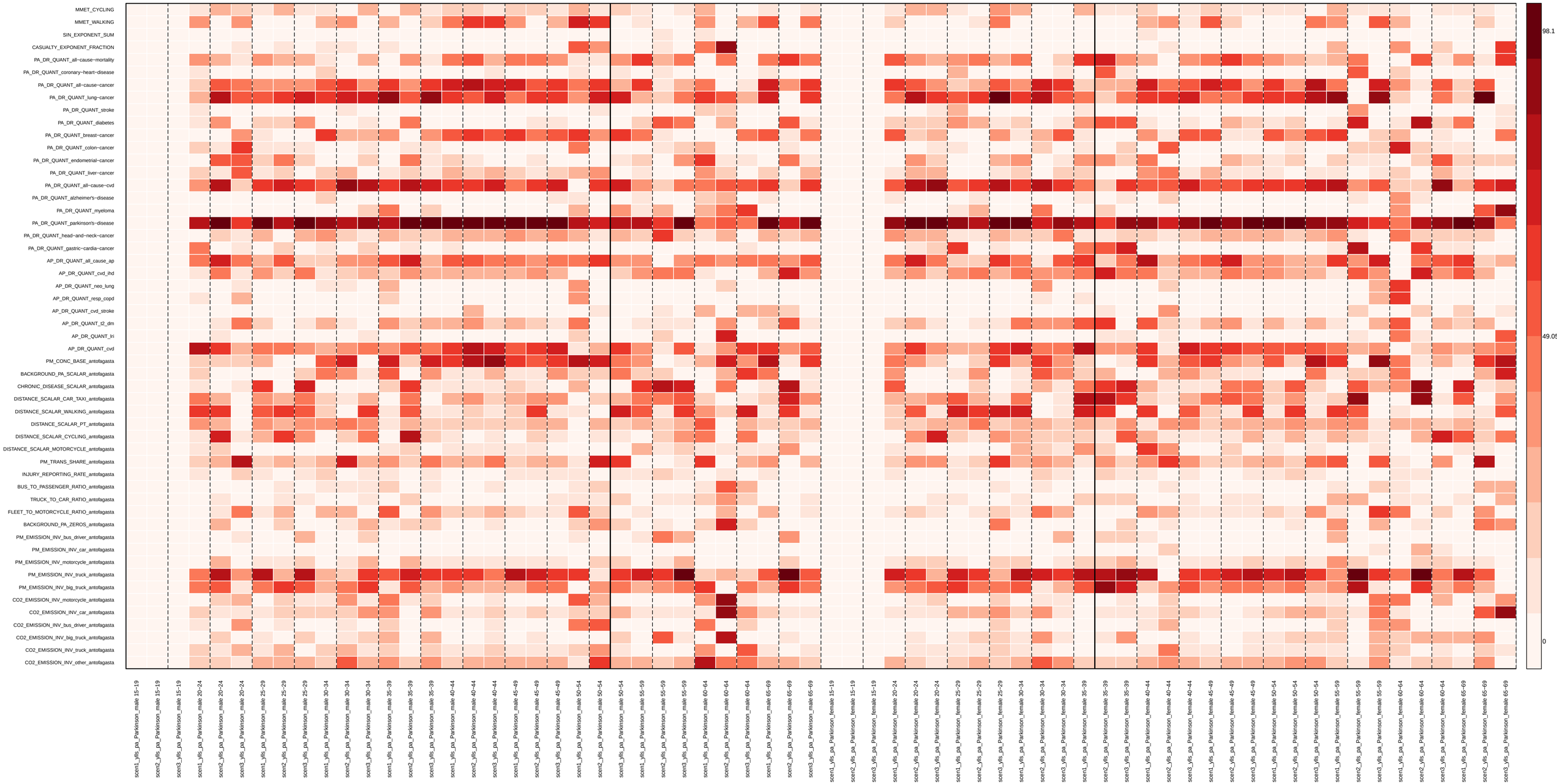
antofagasta − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



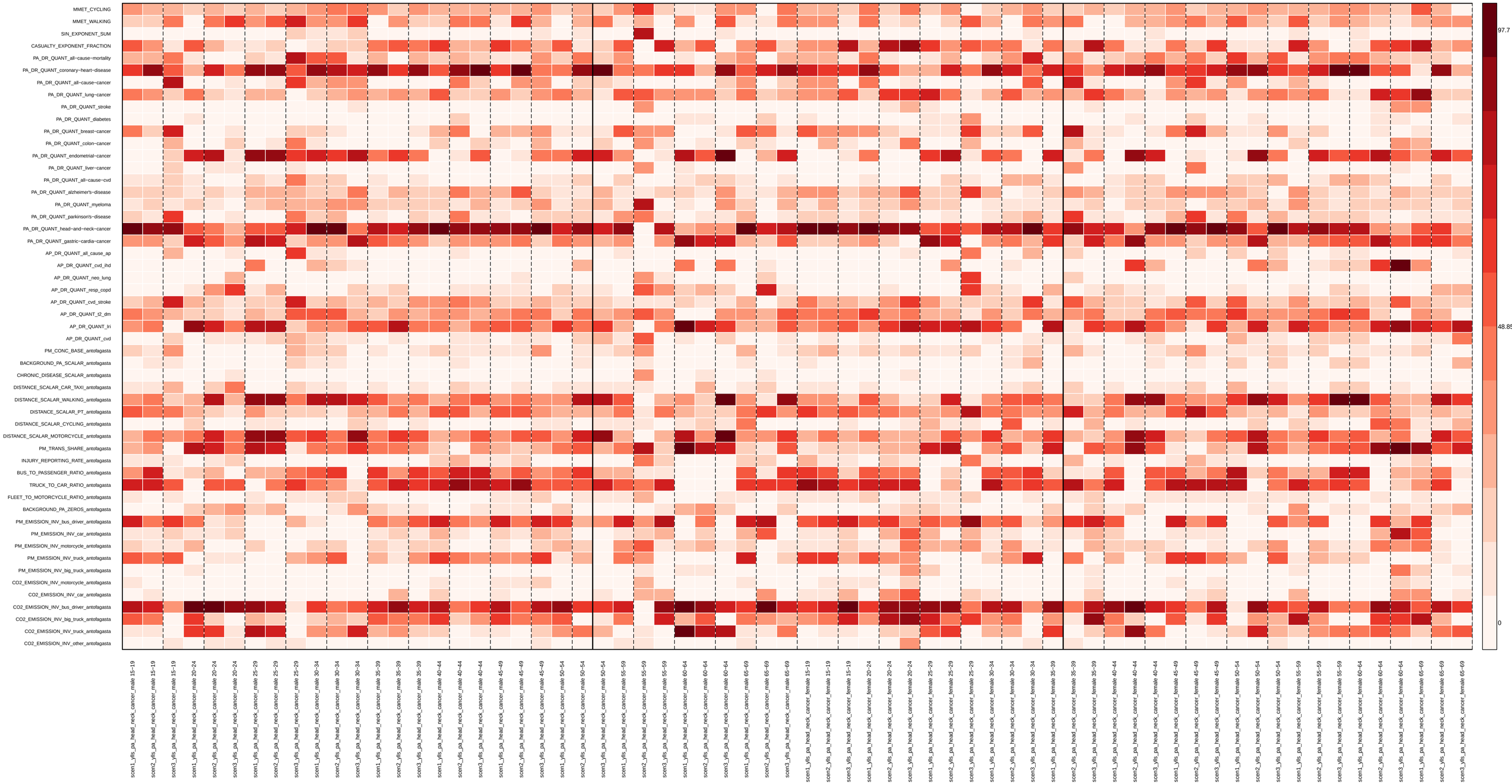
antofagasta − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



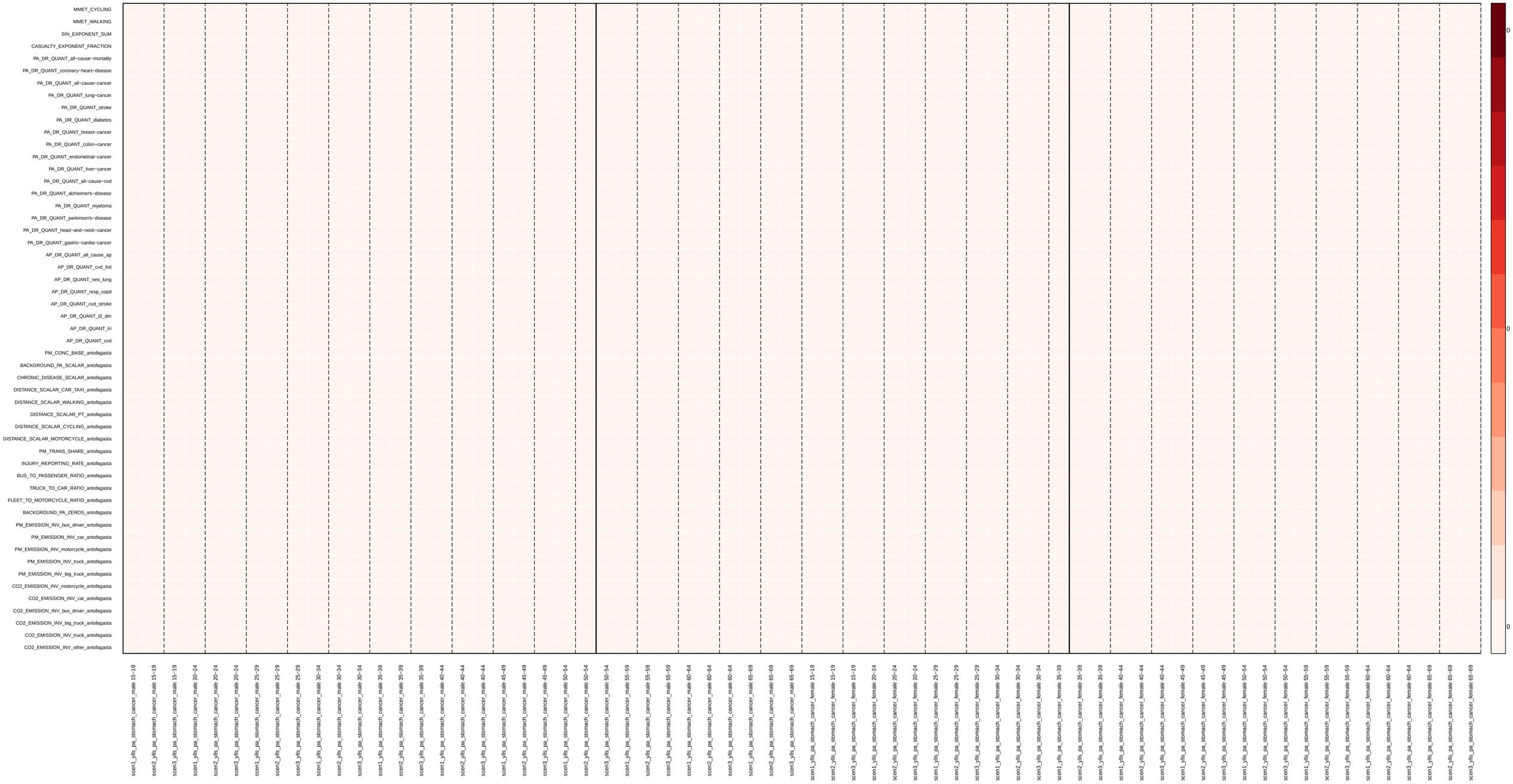
antofagasta − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



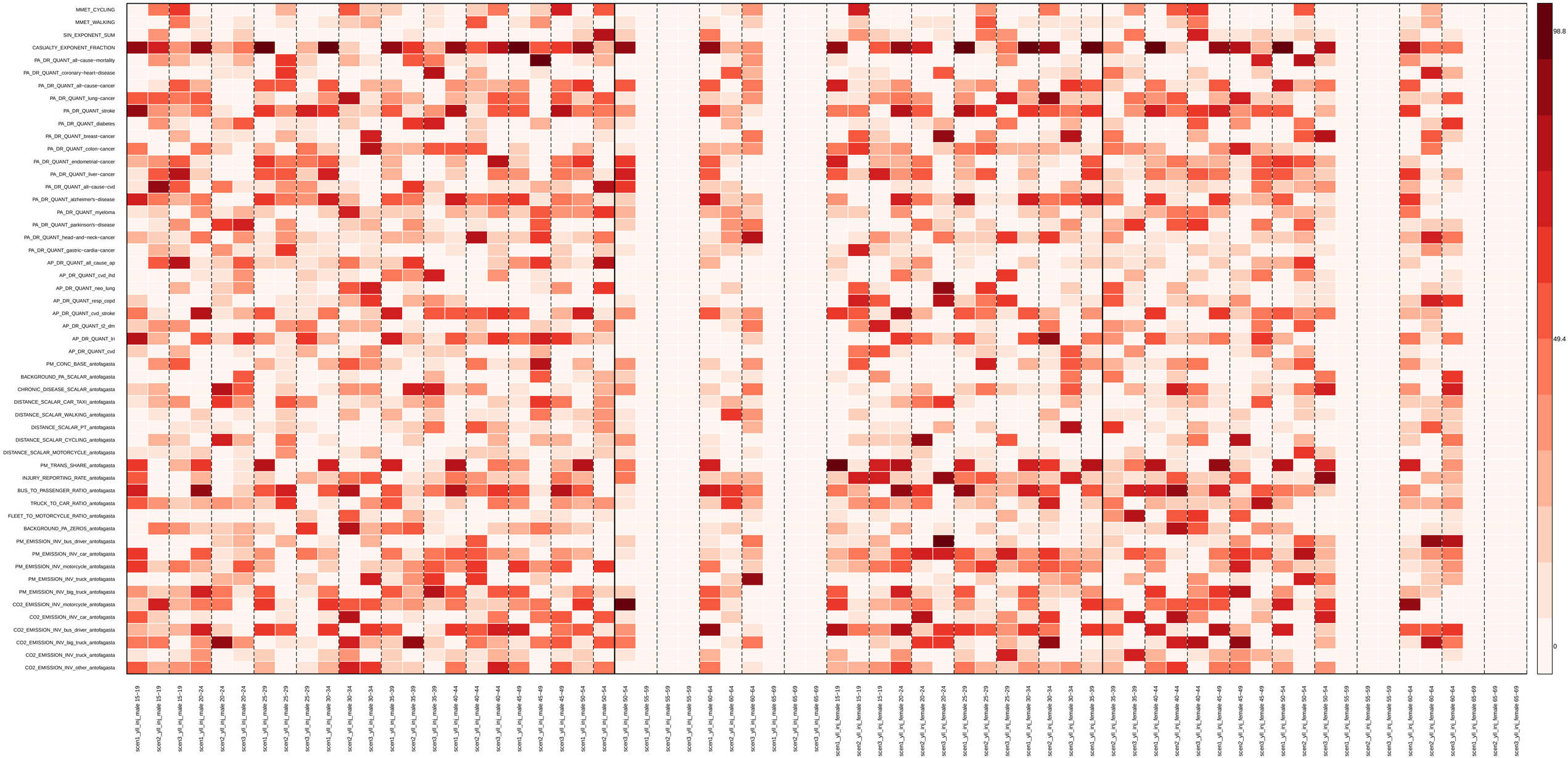
antofagasta − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



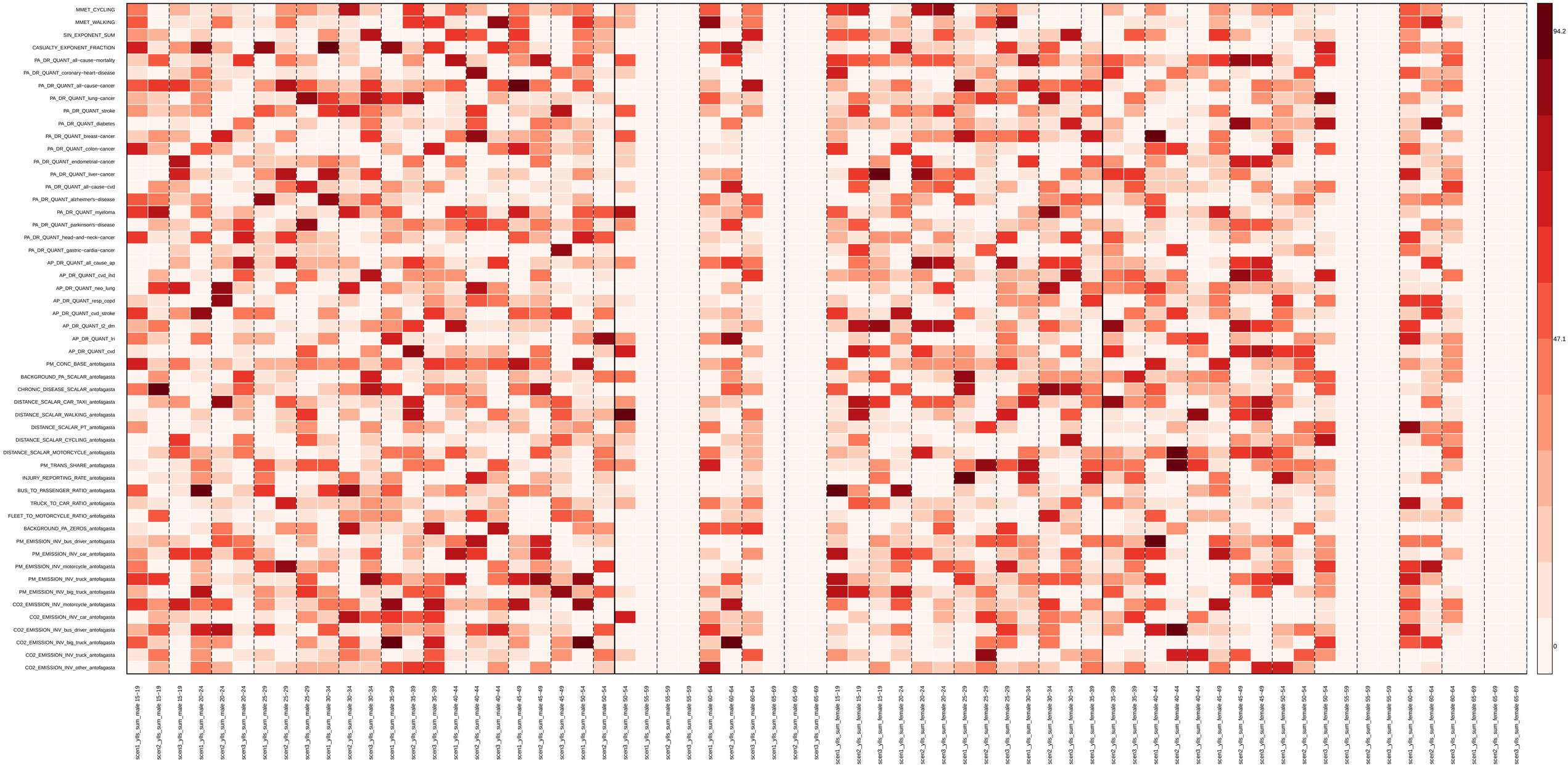
antofagasta − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



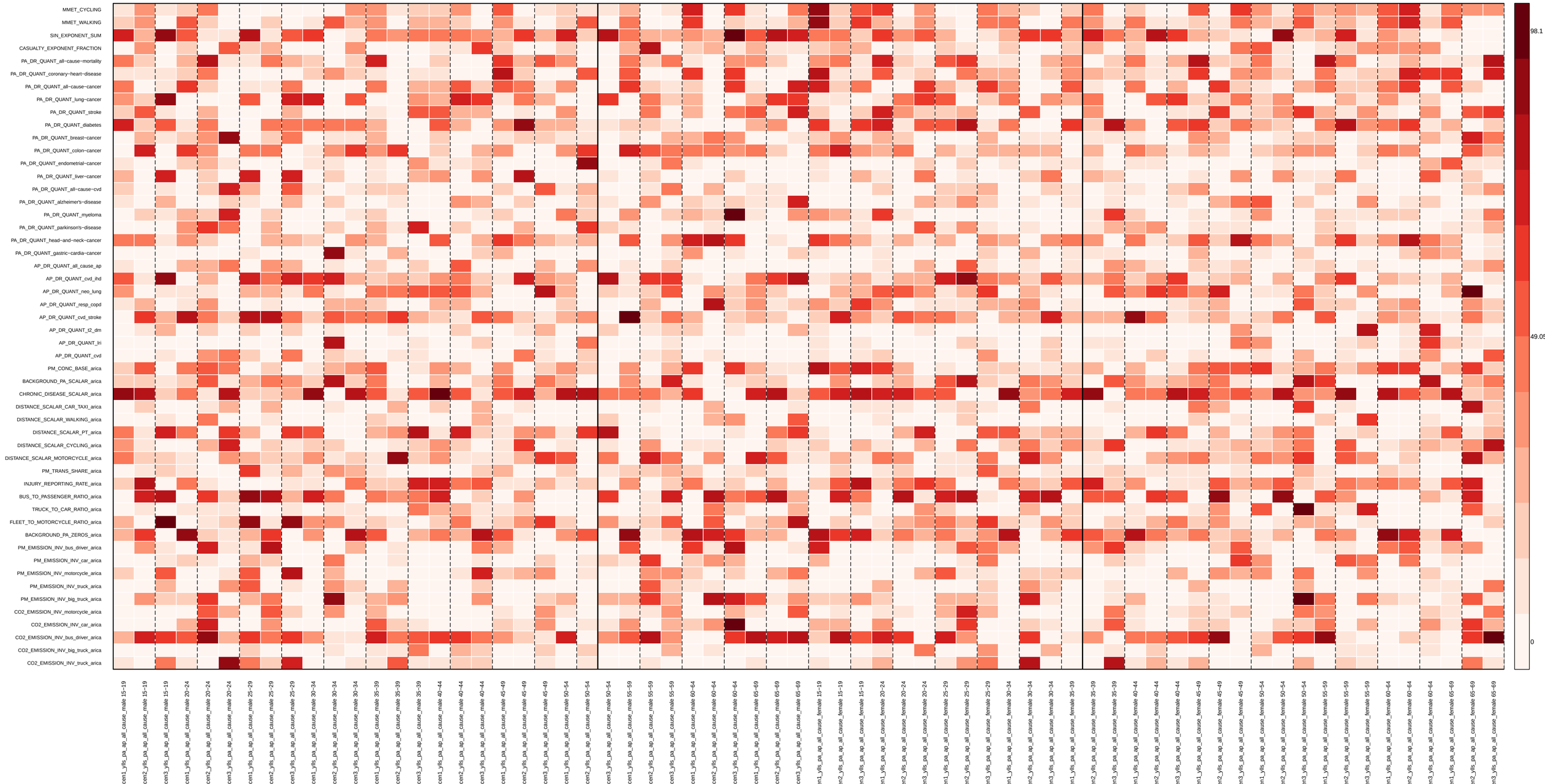
antofagasta − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



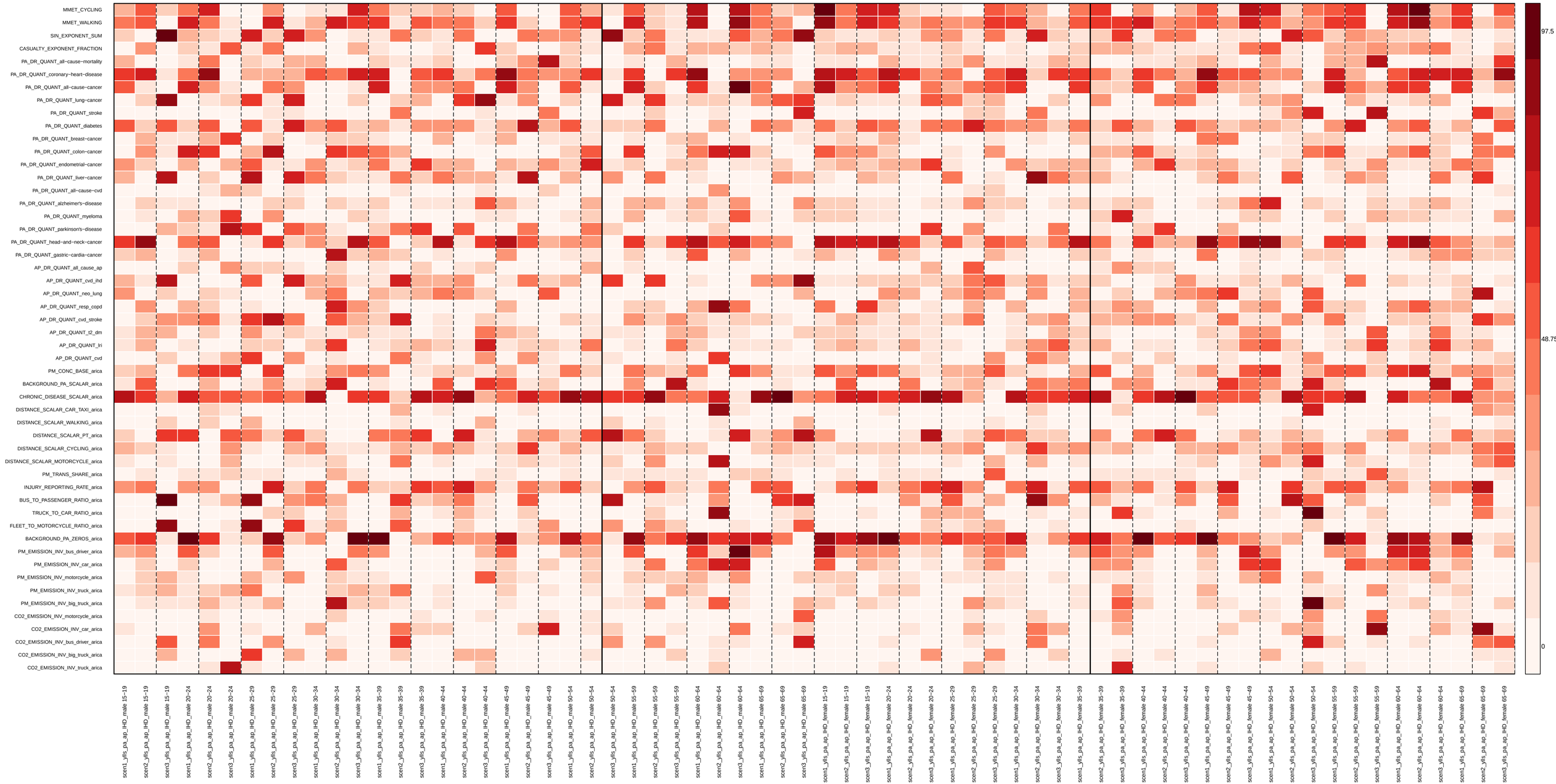
antofagasta − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



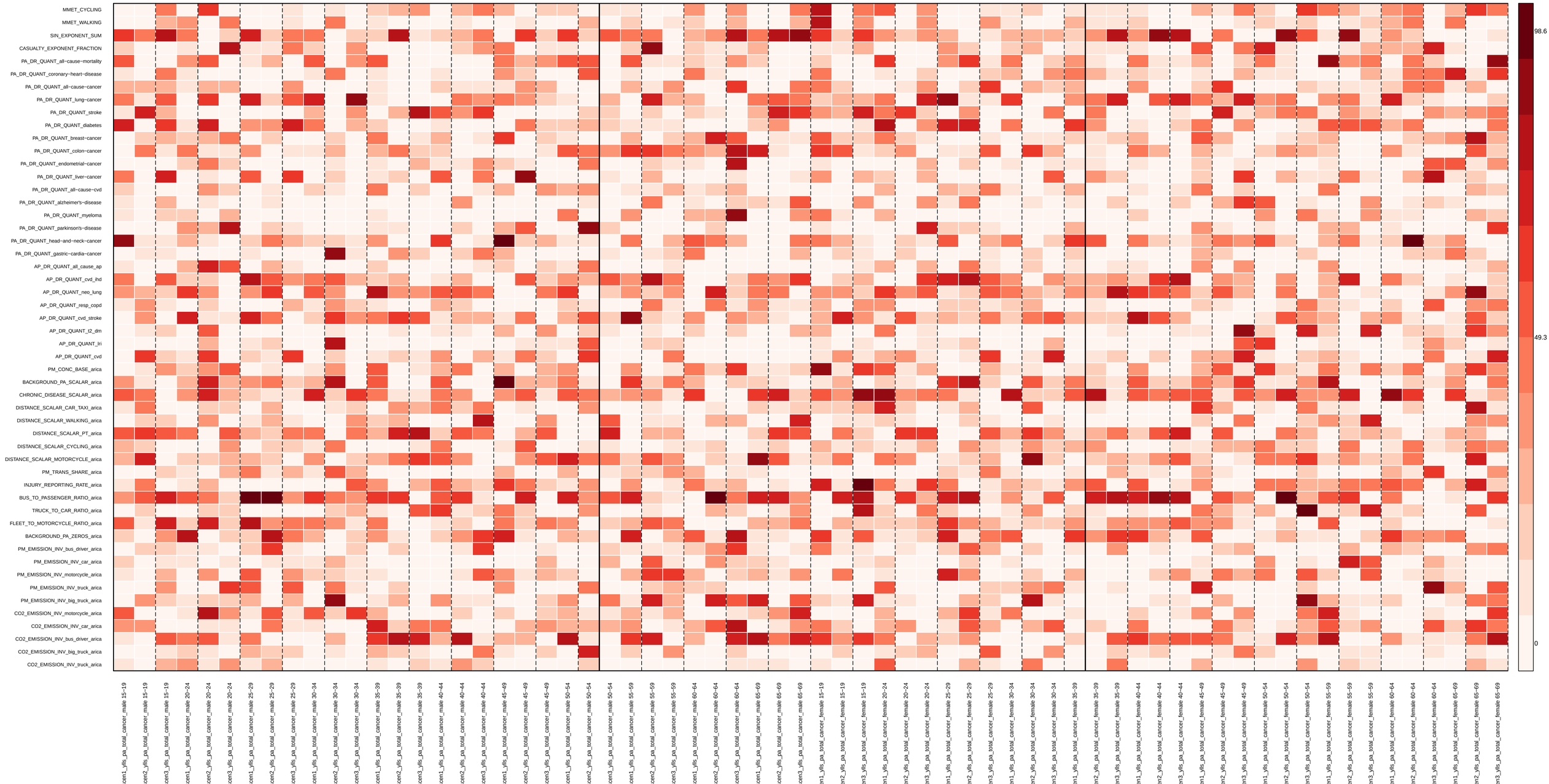
arica − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



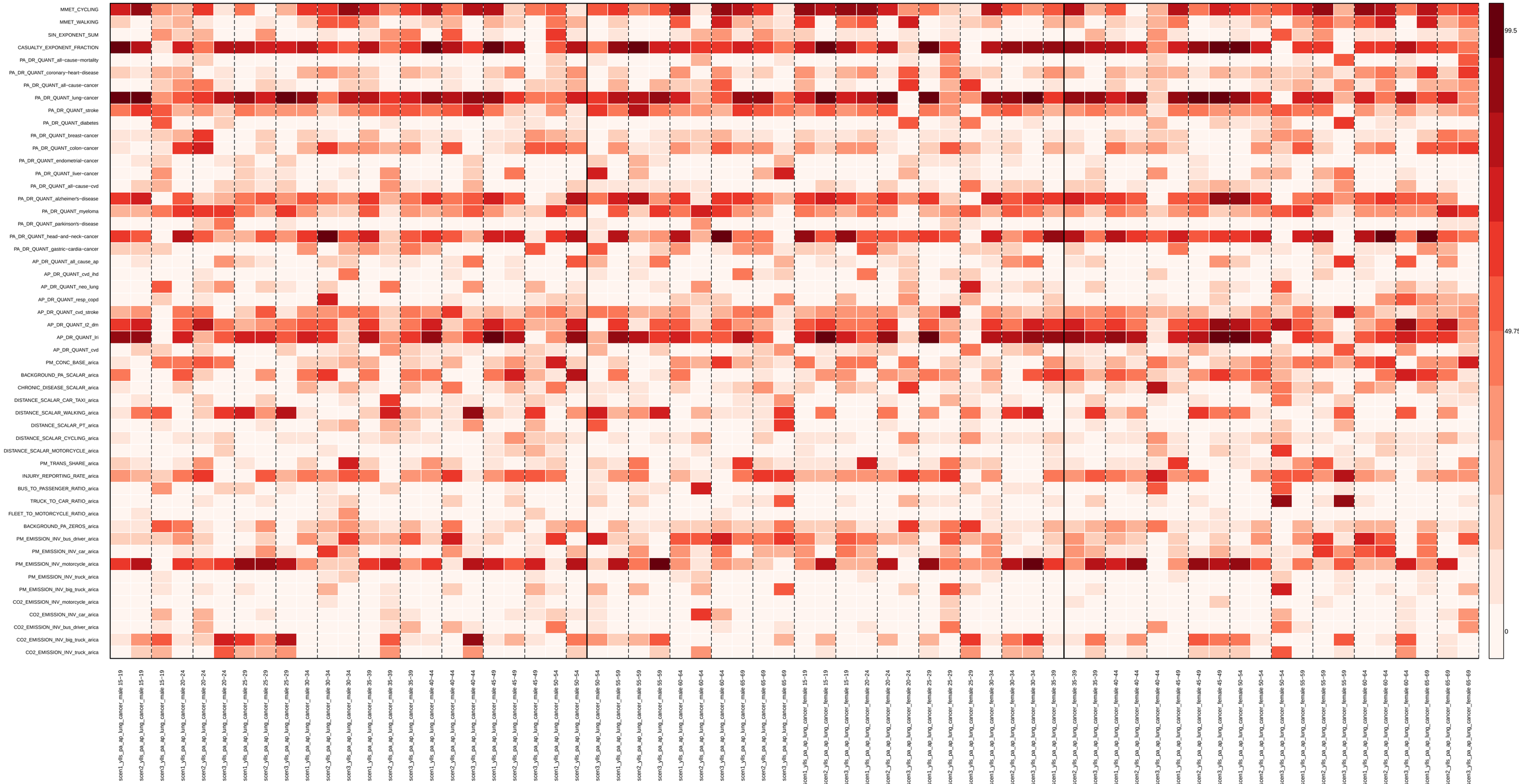
arica − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



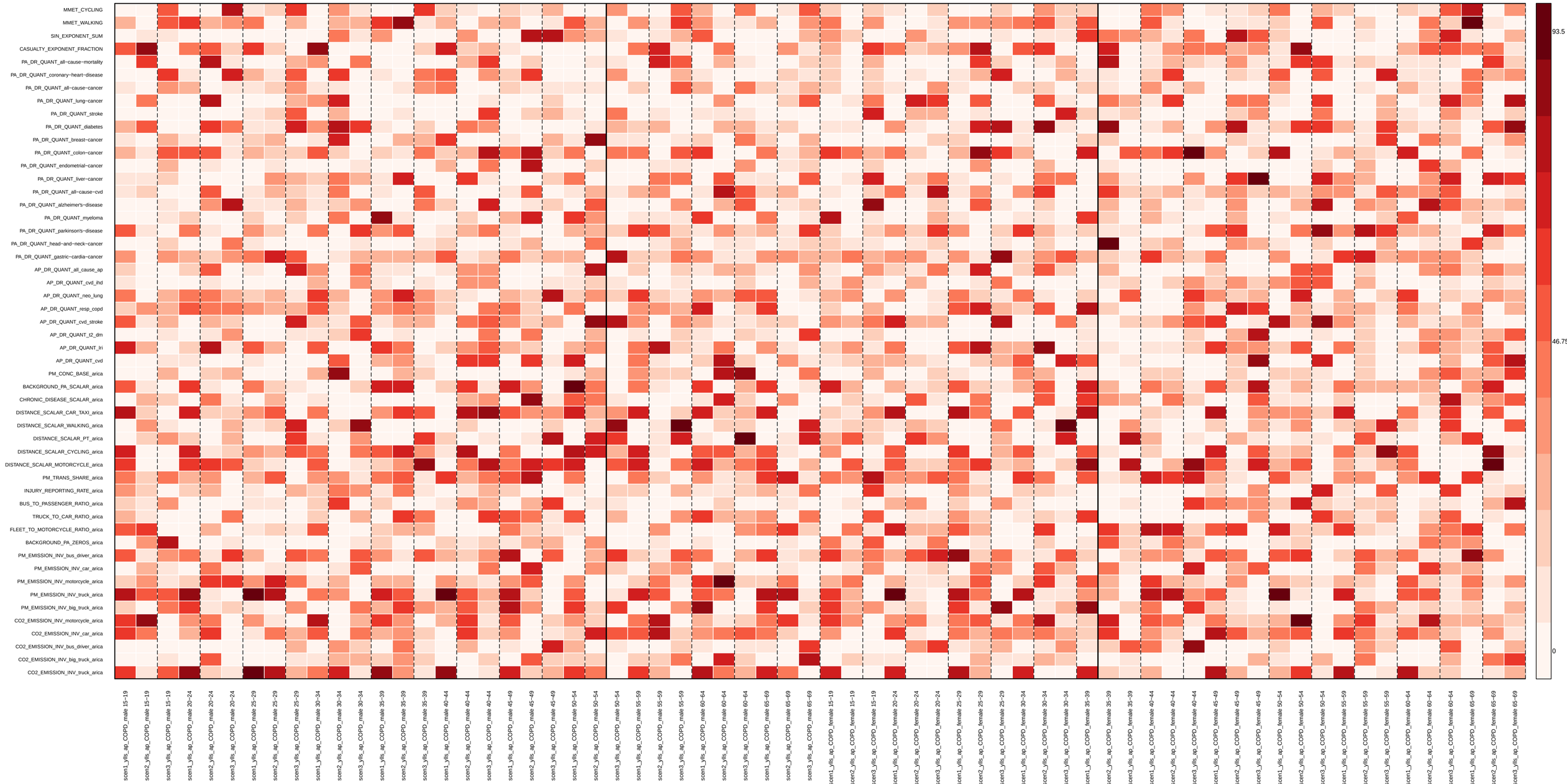
arica − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



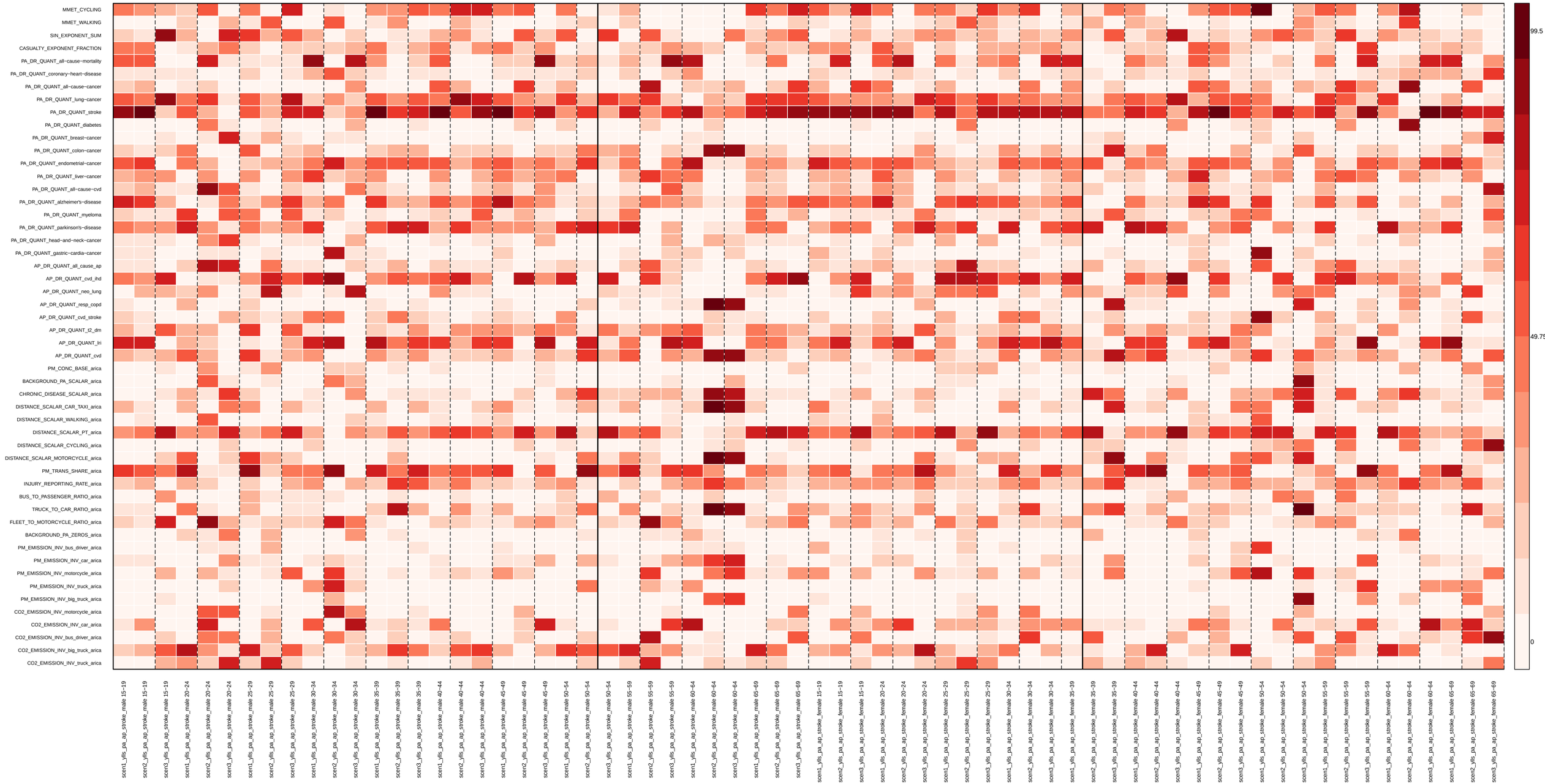
arica − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



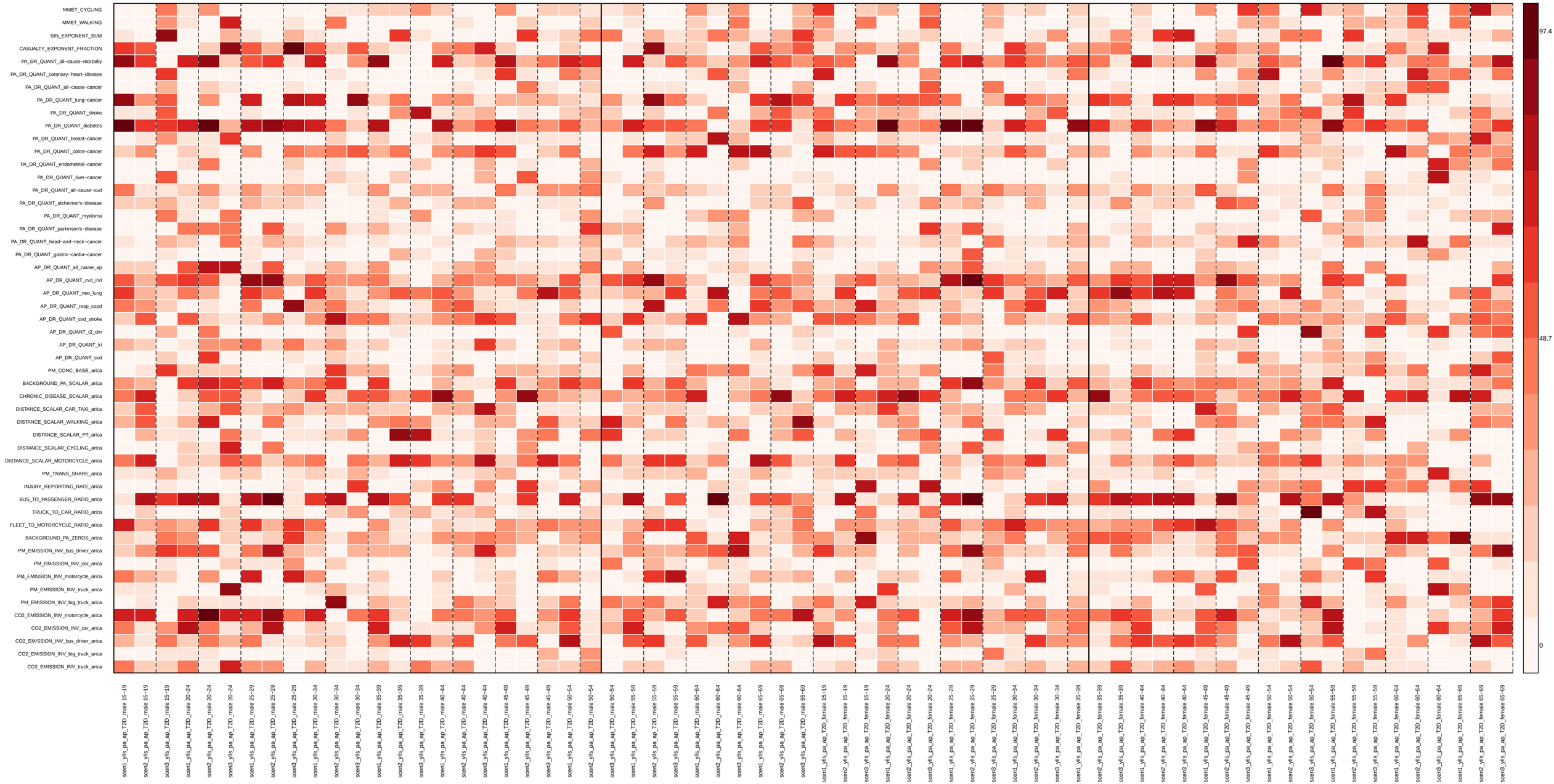
arica − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



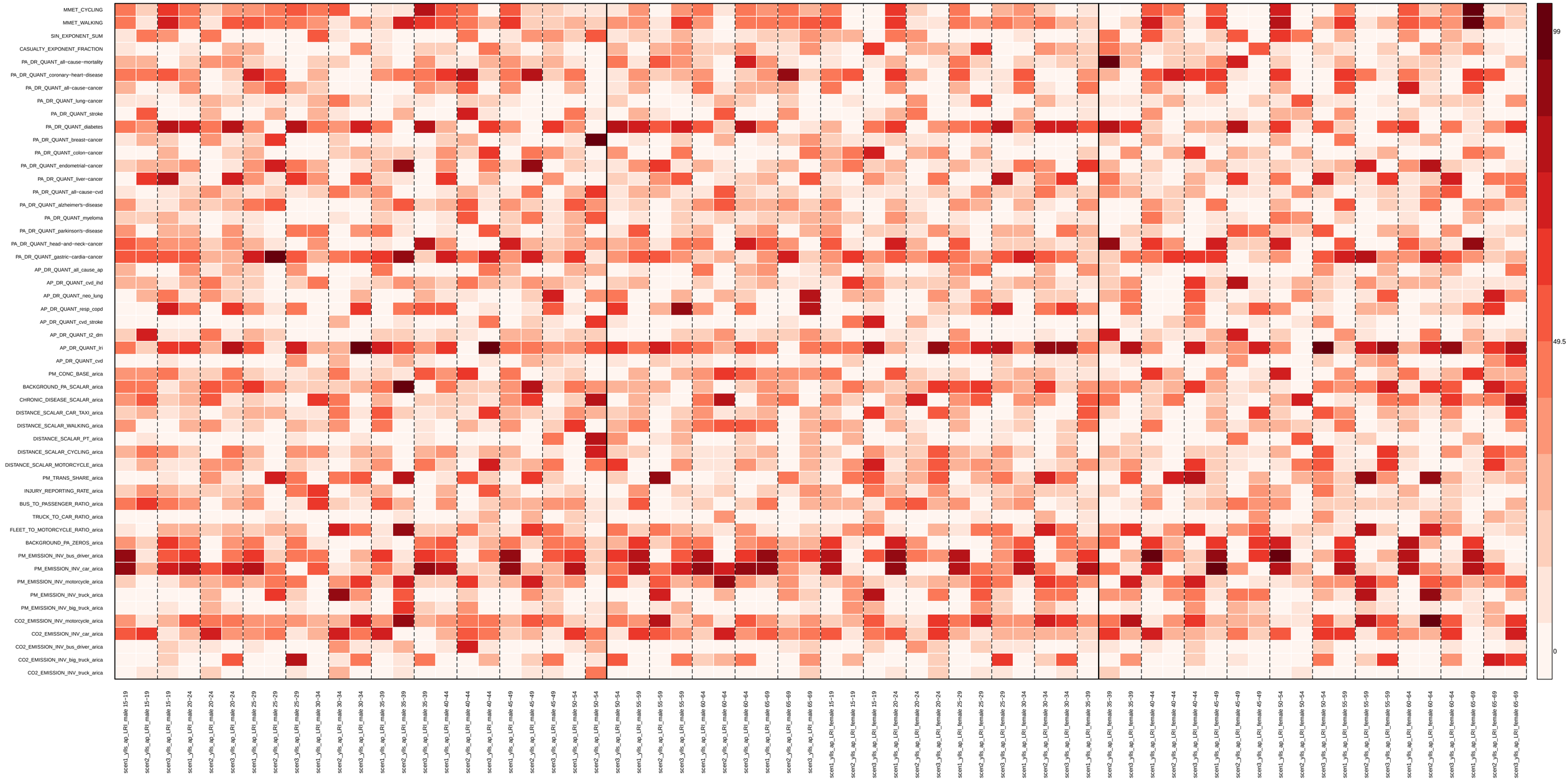
arica − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



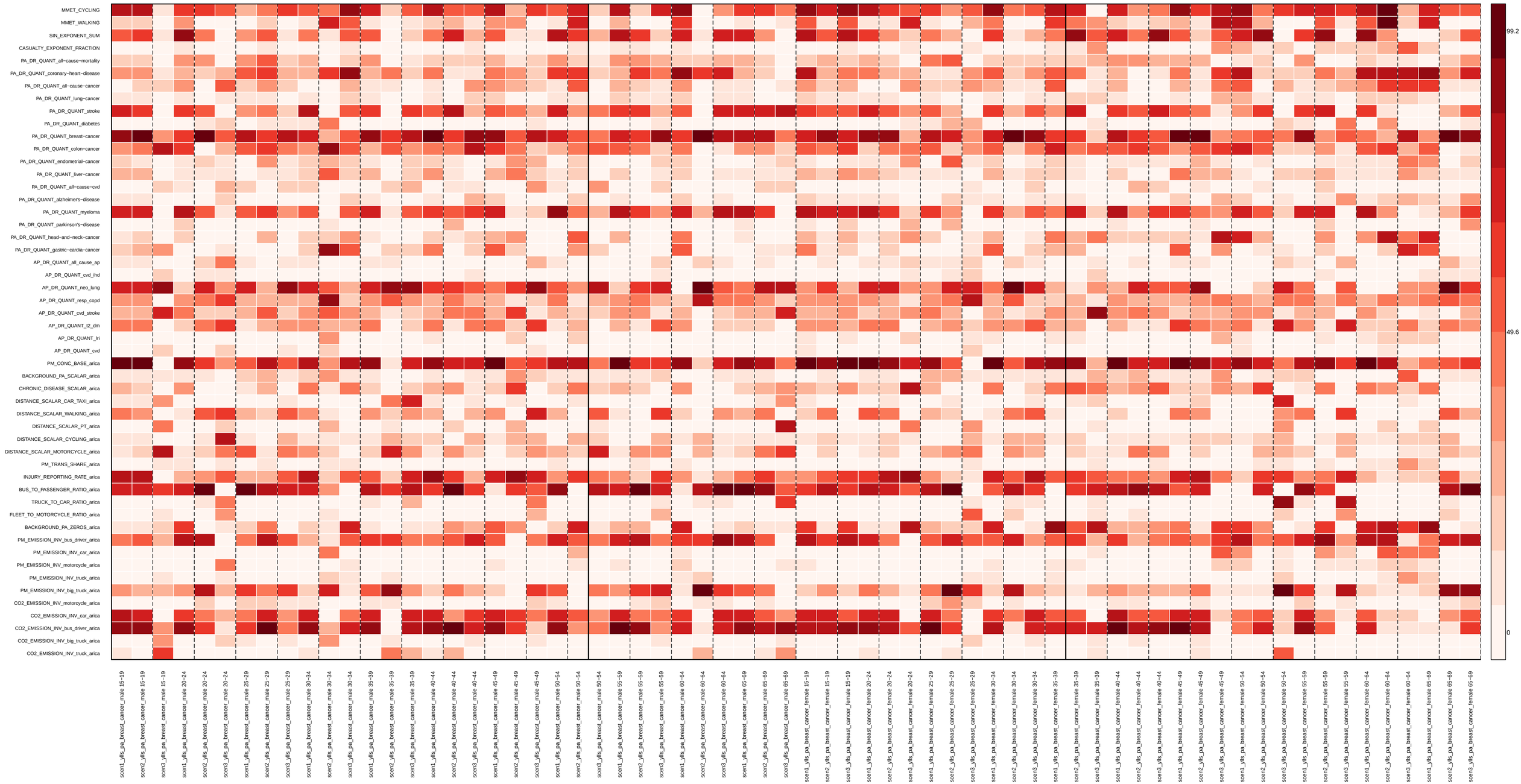
arica − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



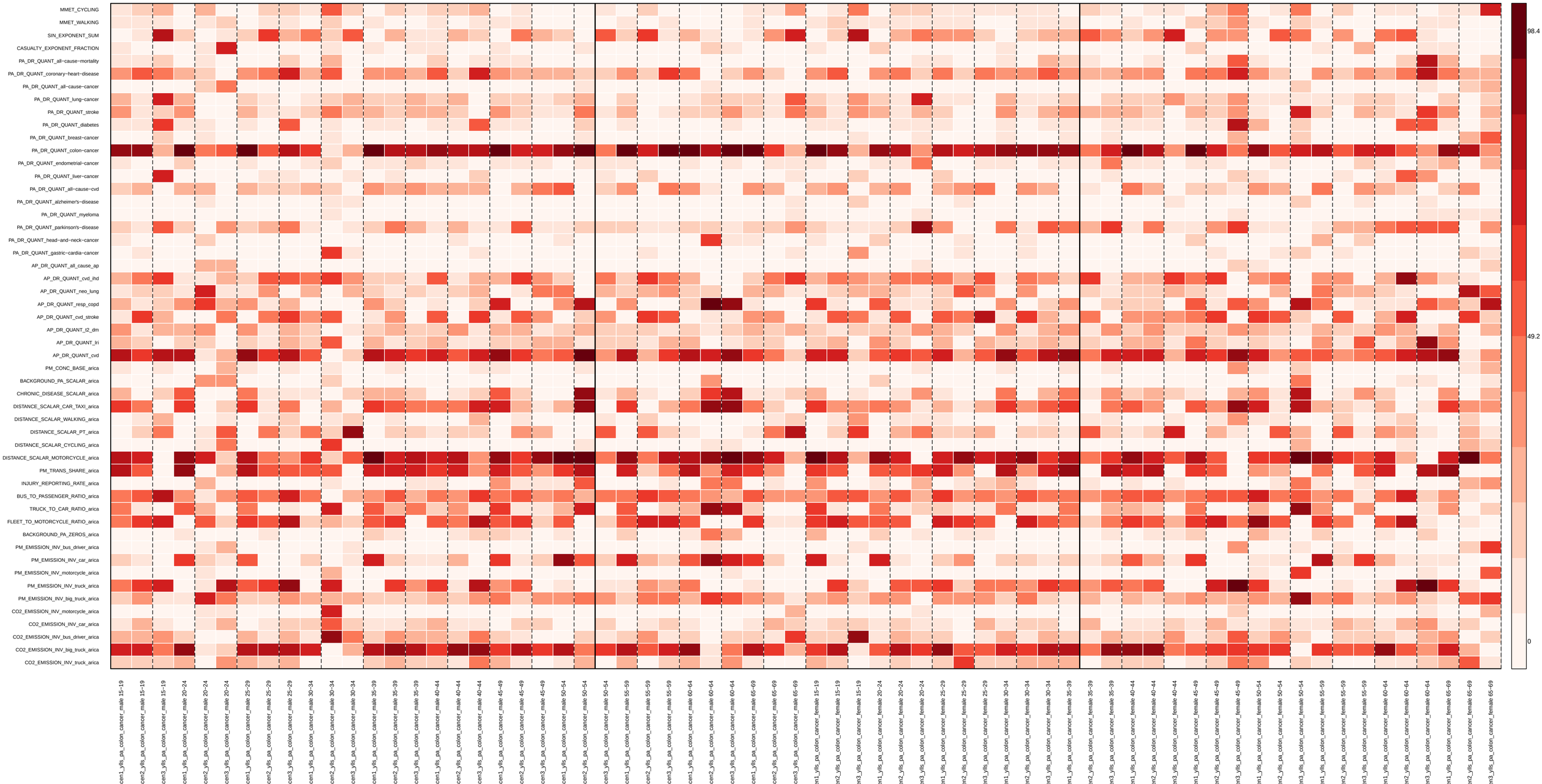
arica − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



arica − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

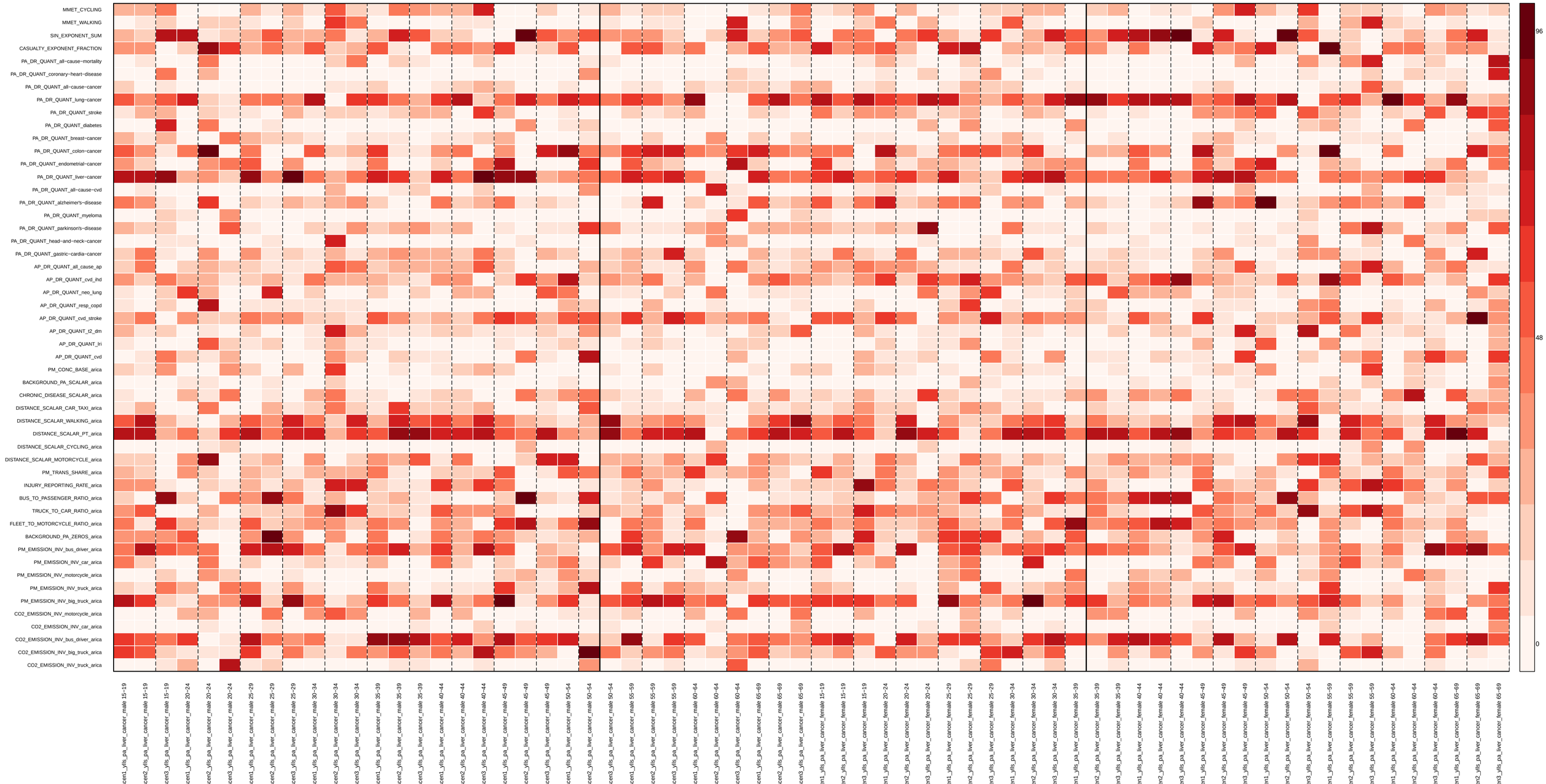


arica − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

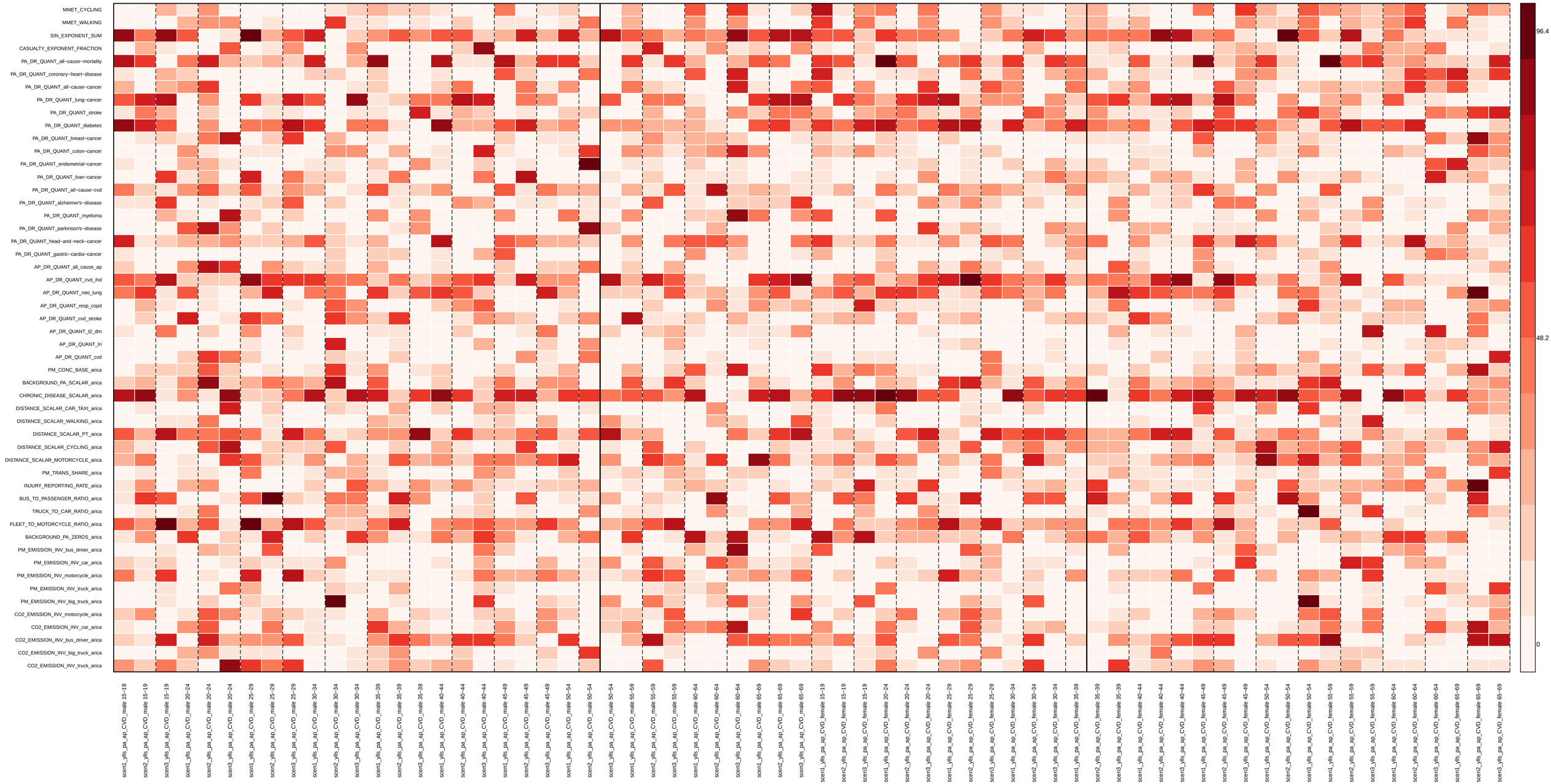


arica − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

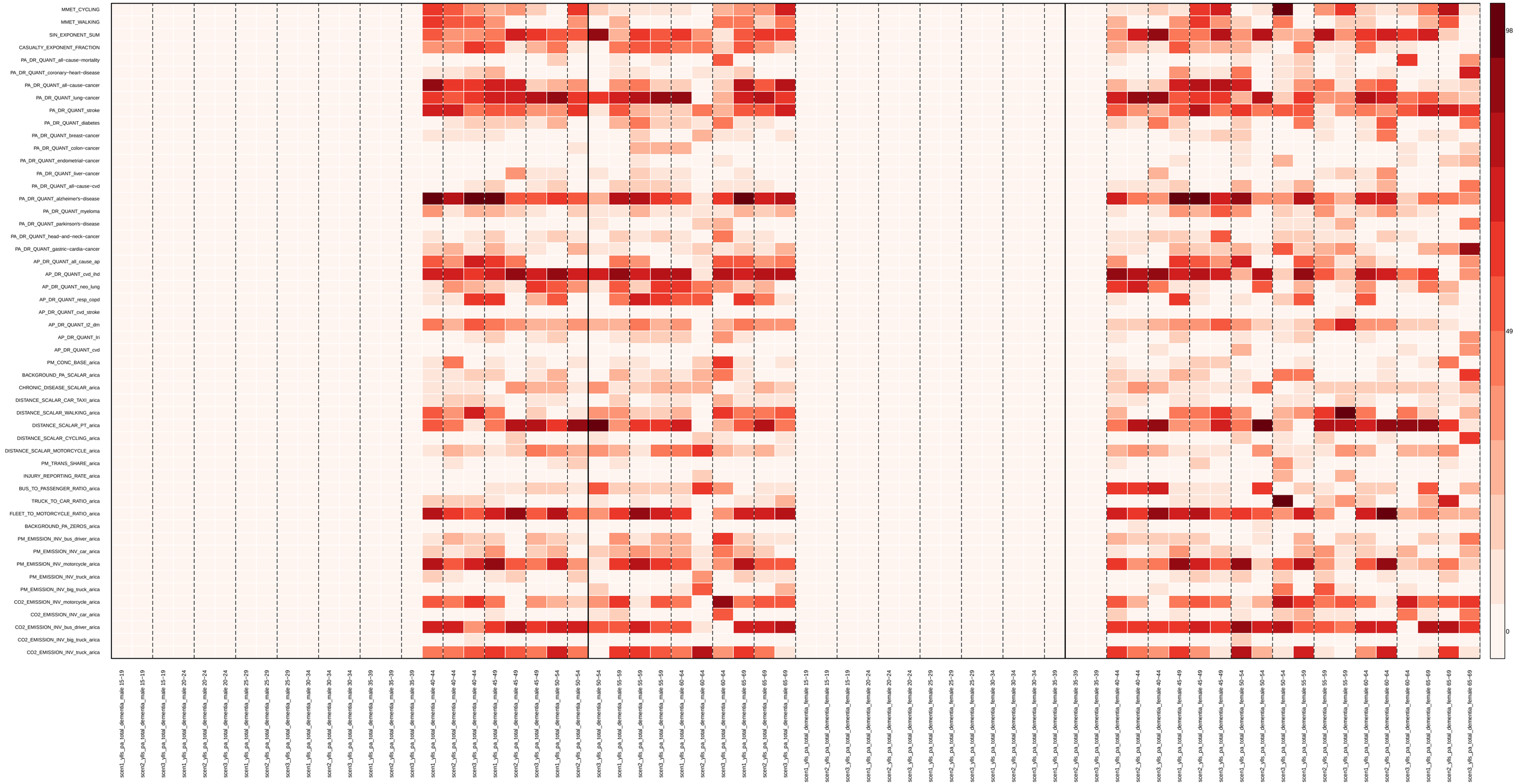
arica – pa_liver_cancer , No of samples: 10 –
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



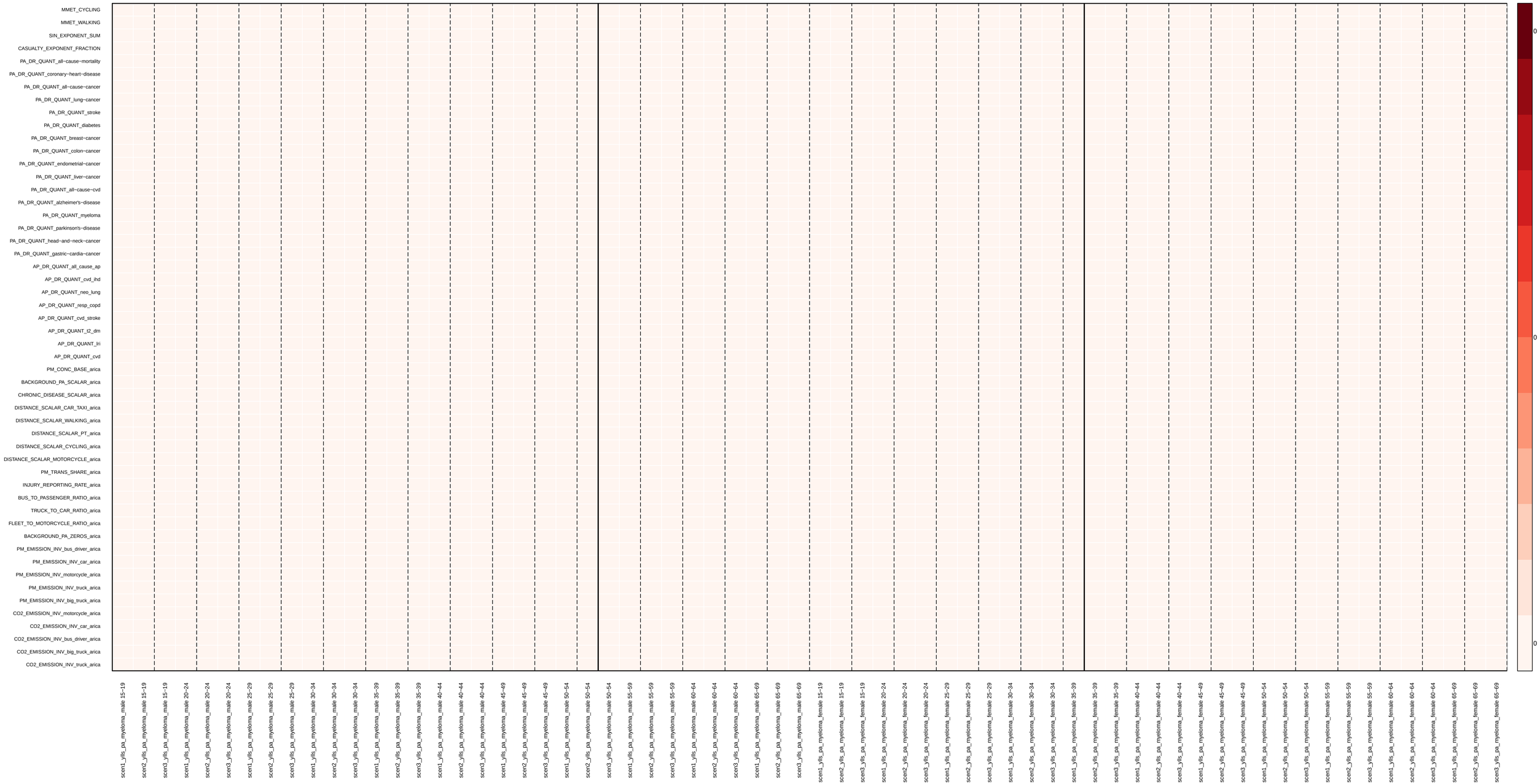
arica − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



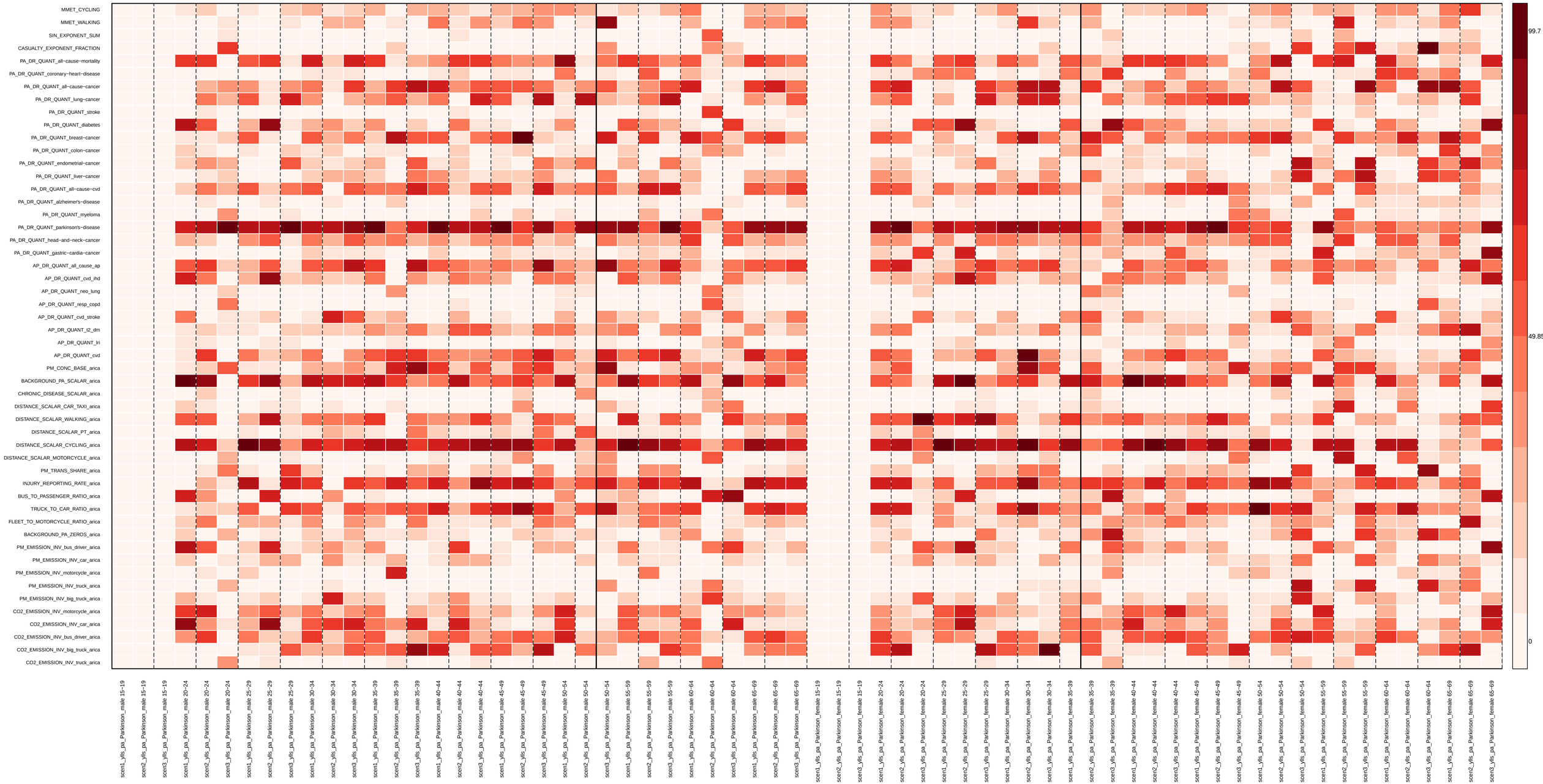
arica − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



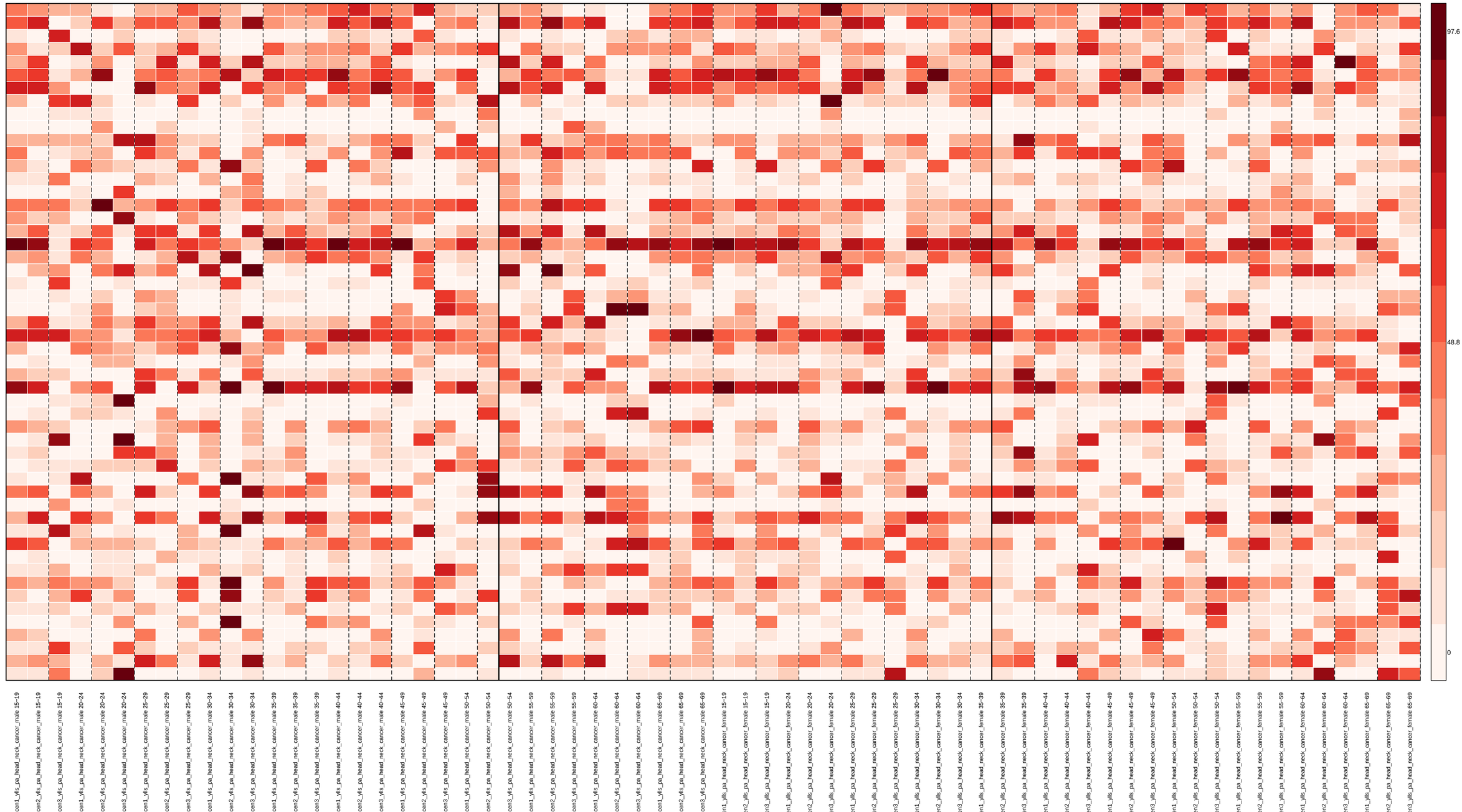
arica − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



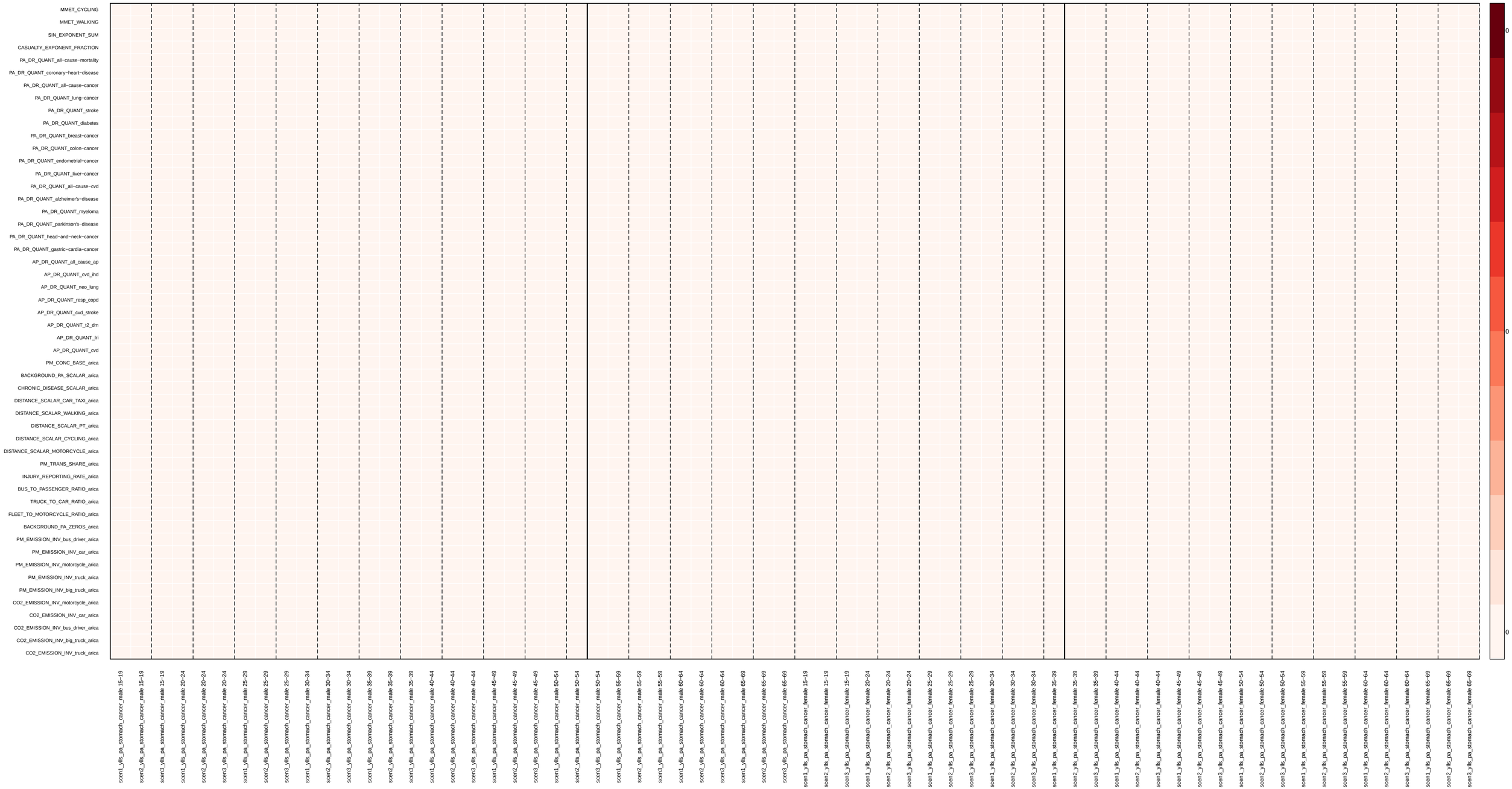
arica − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



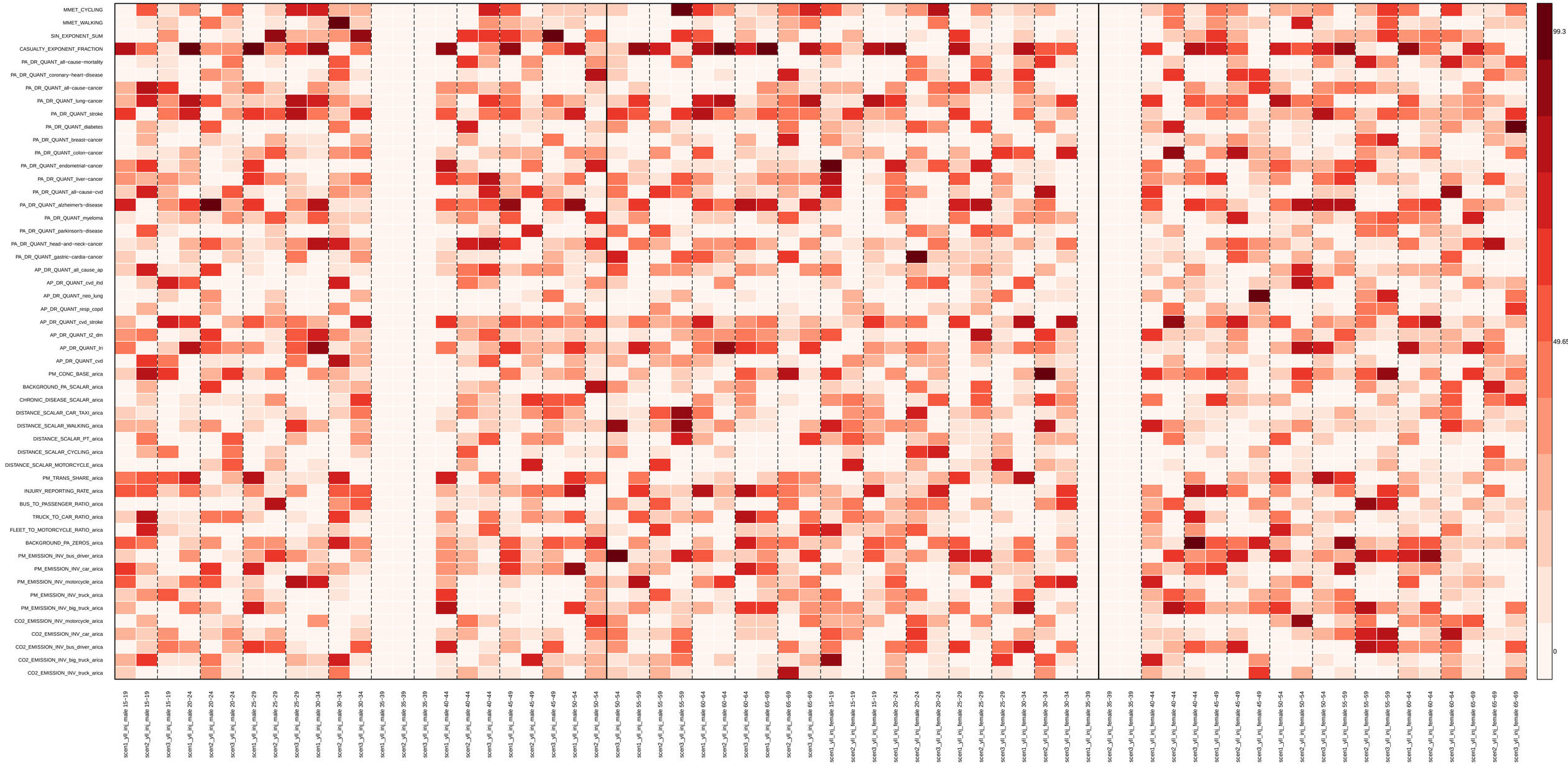
arica − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



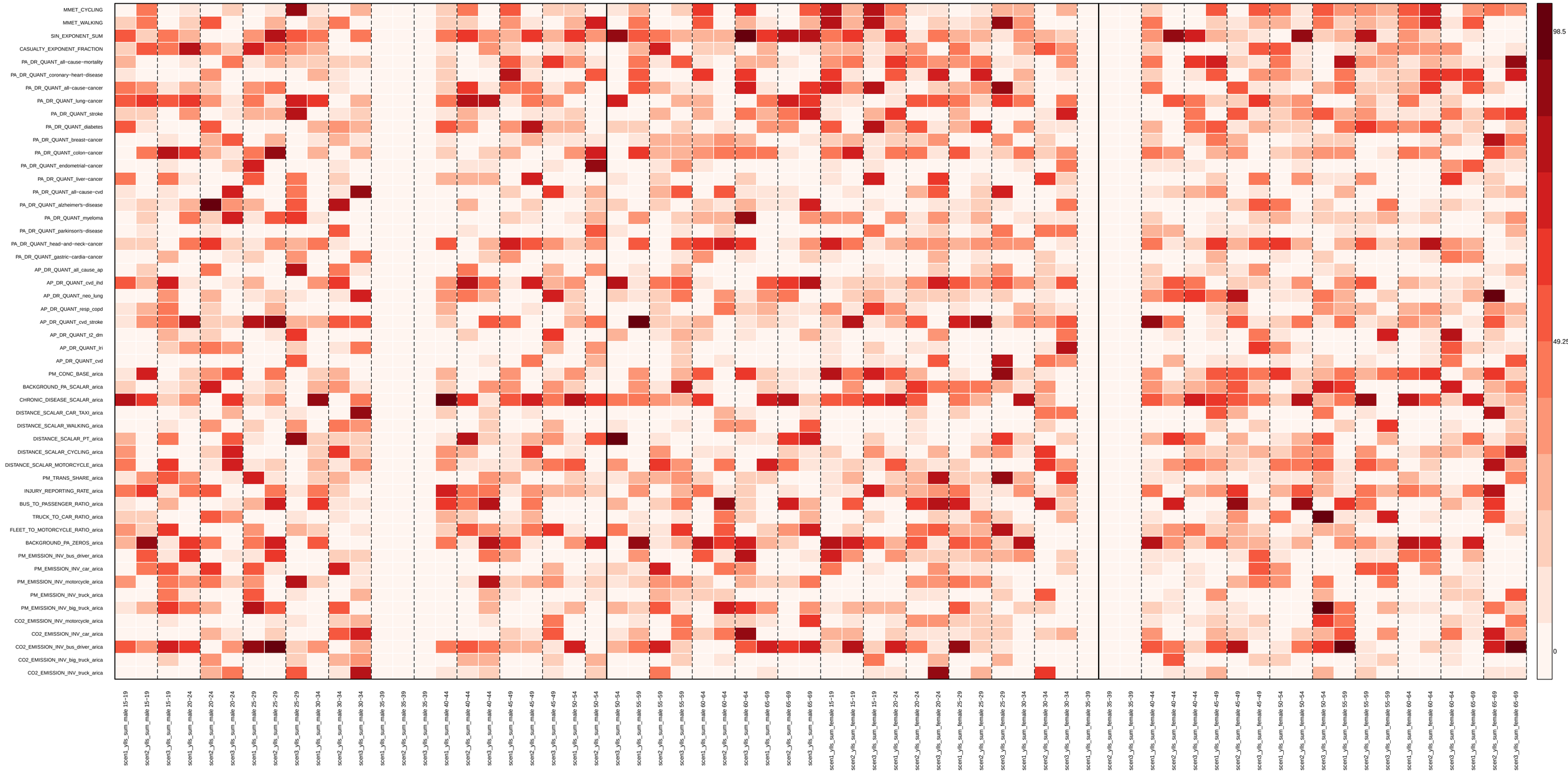
arica − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



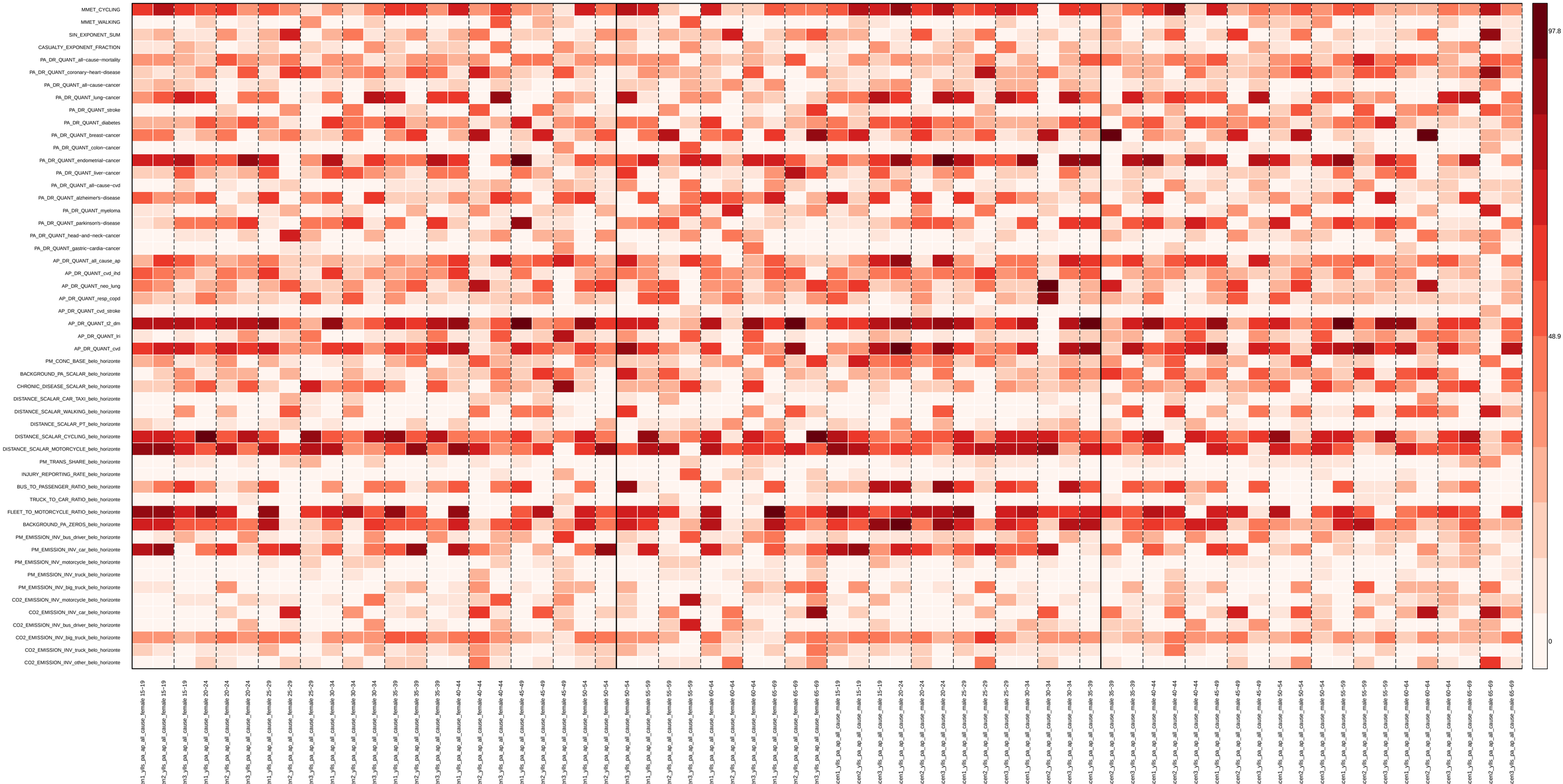
arica − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



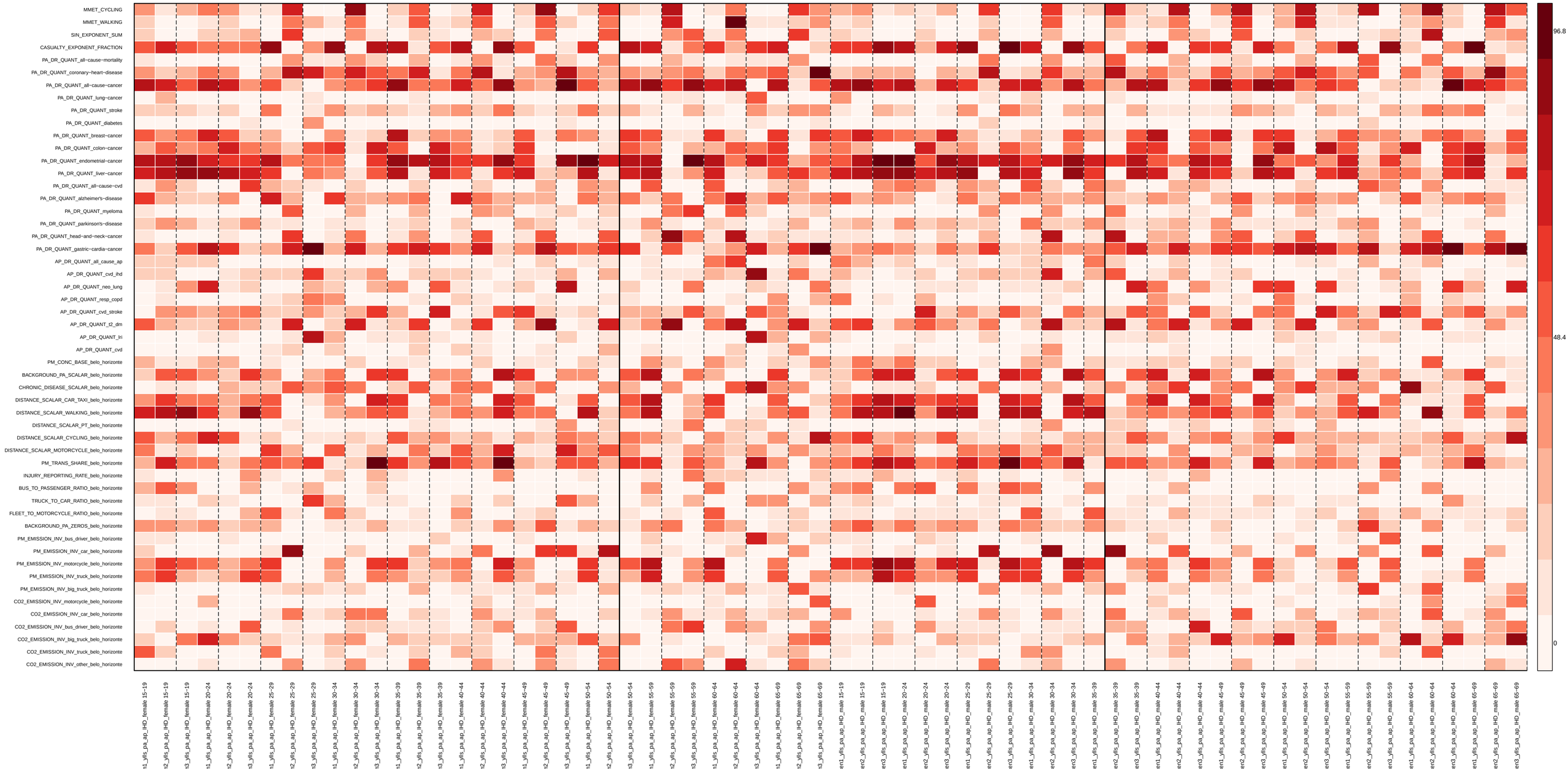
arica − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



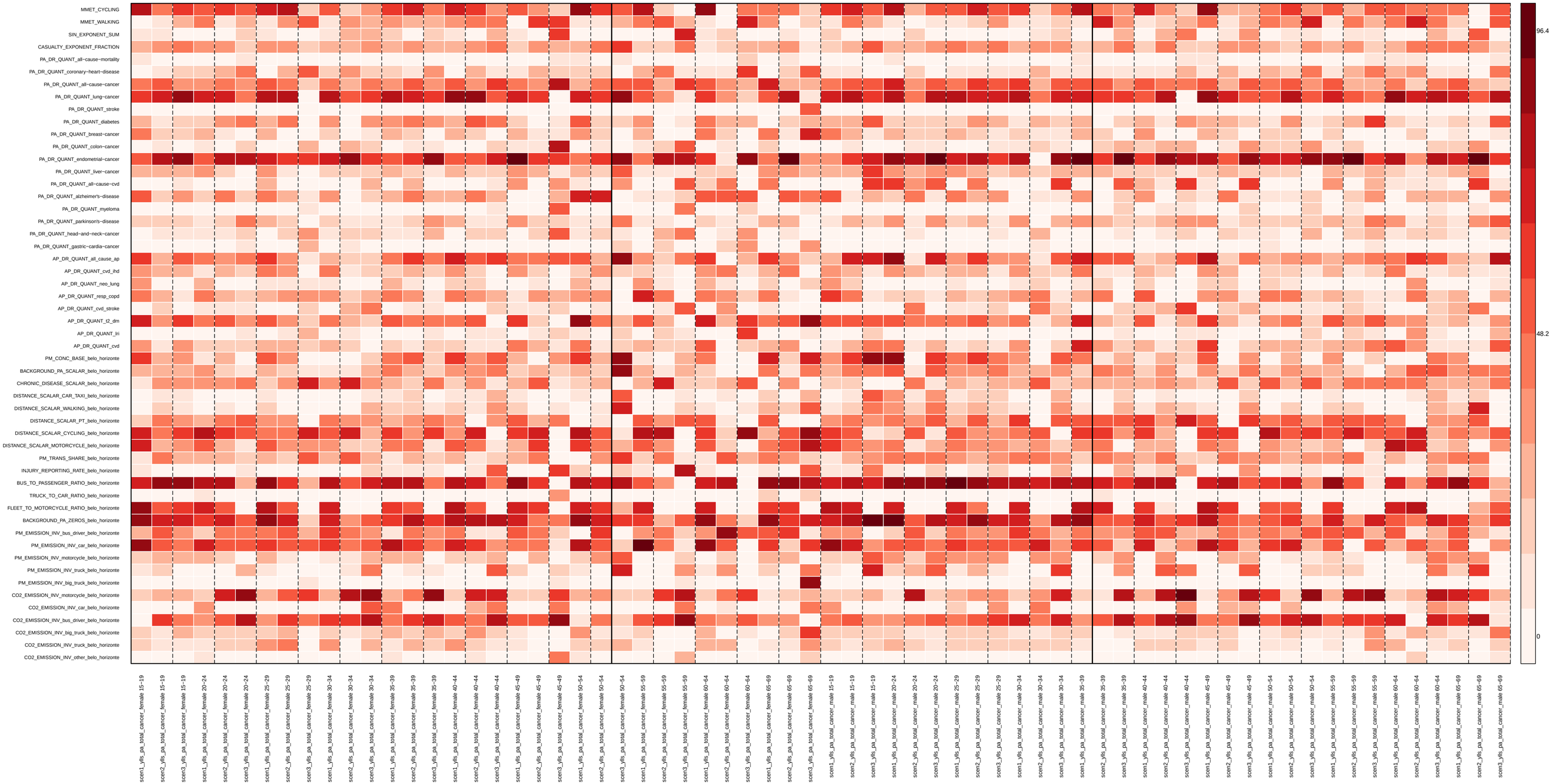
belo_horizonte − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



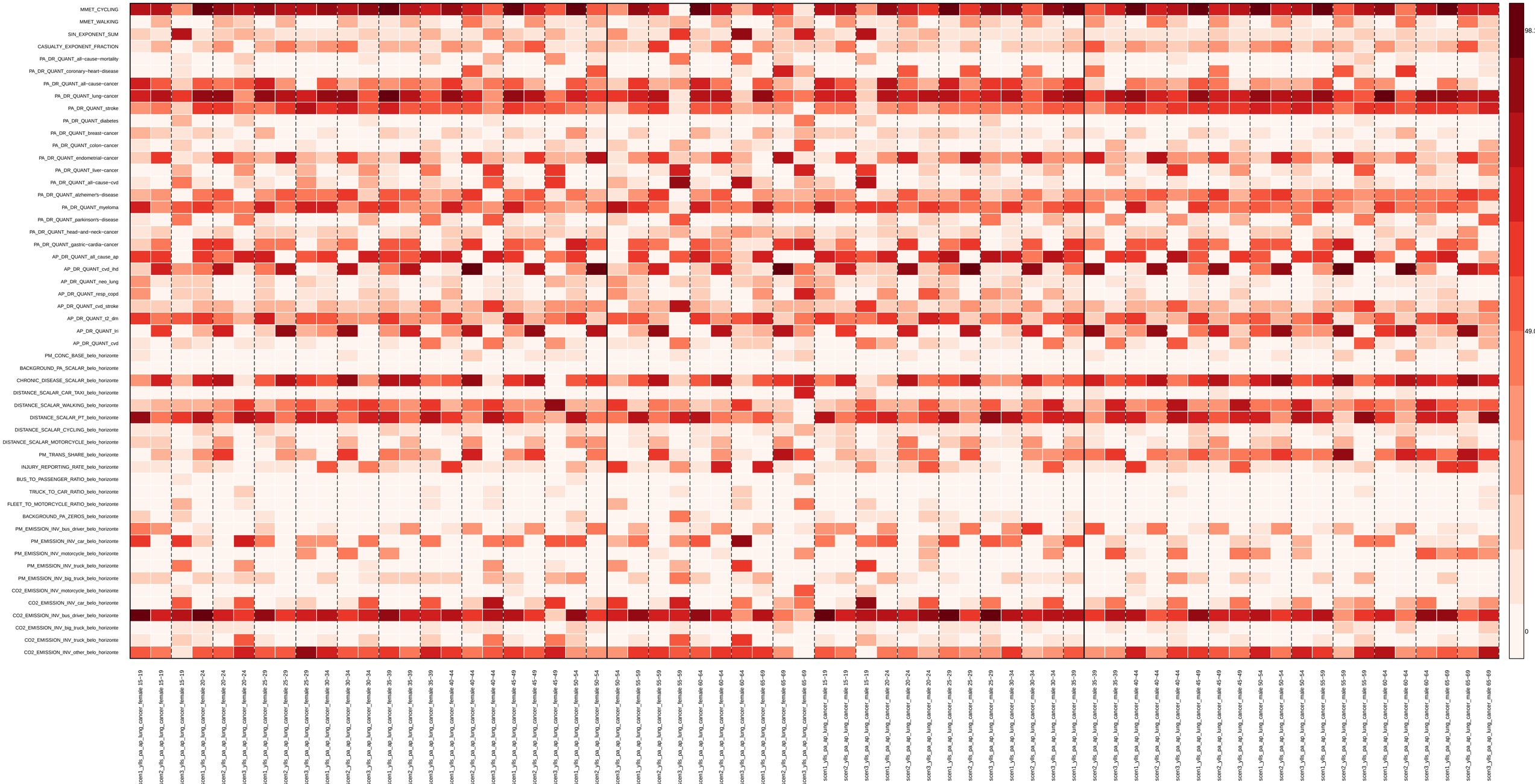
belo_horizonte − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



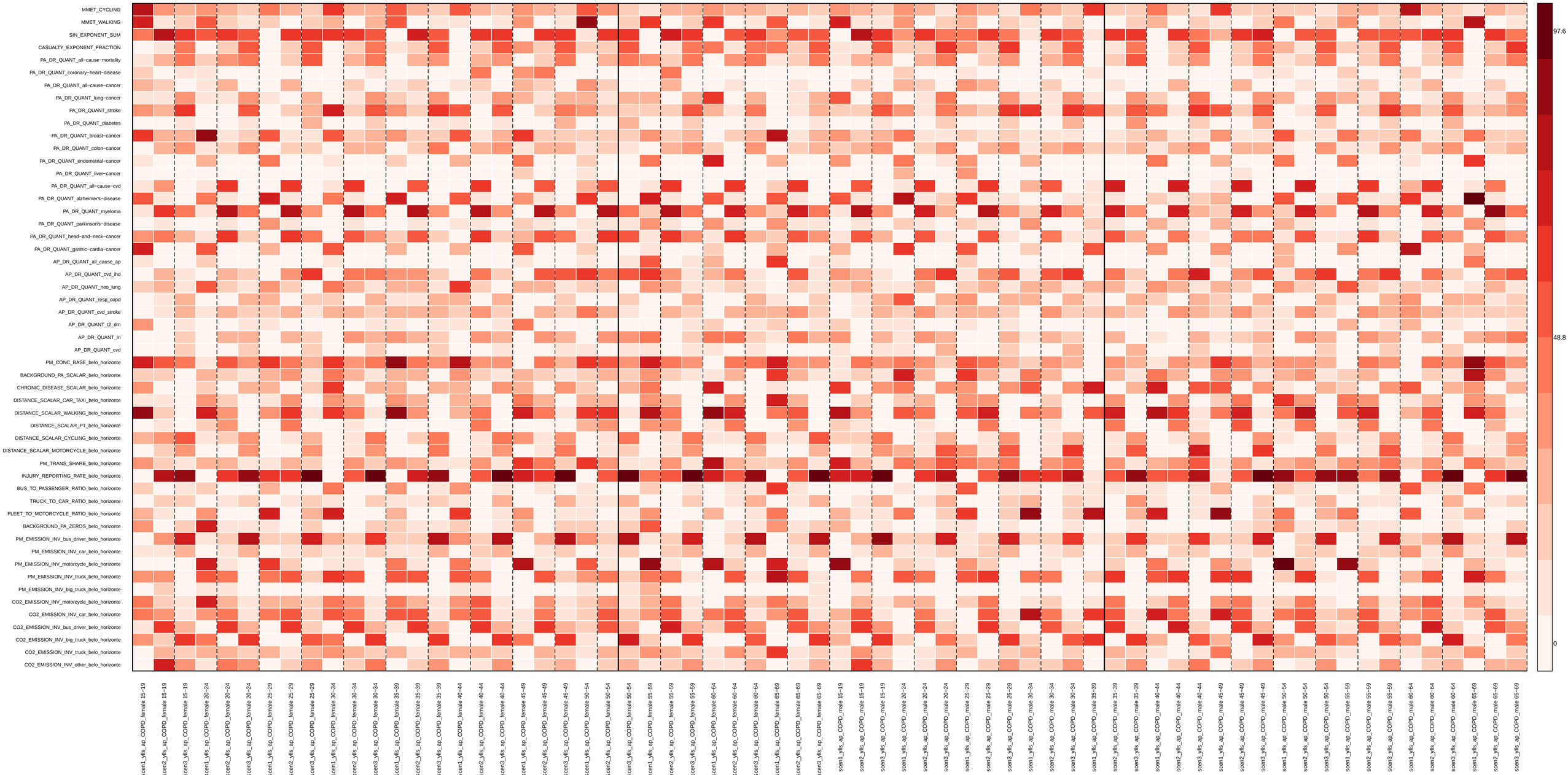
belo_horizonte − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



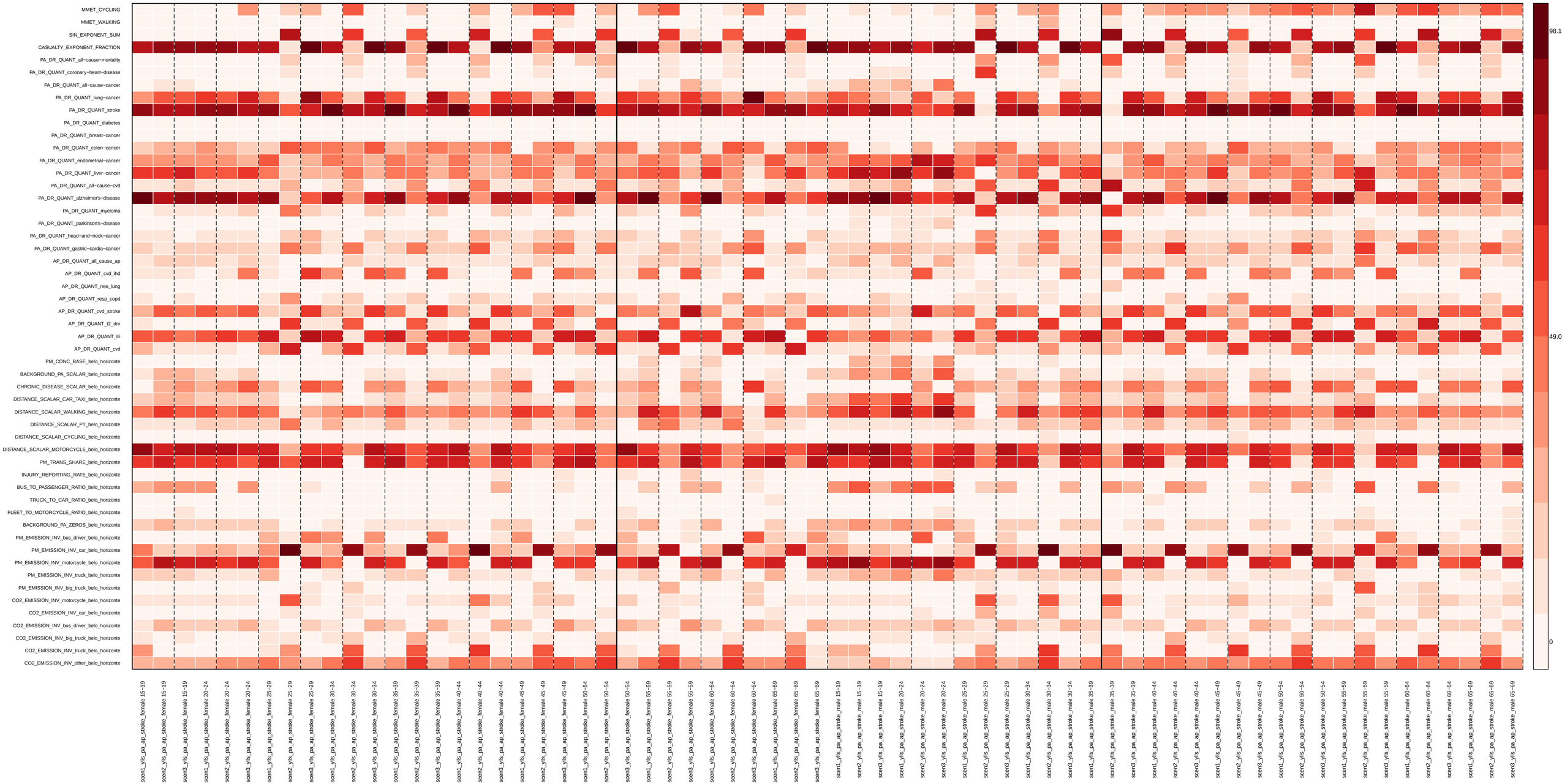
belo_horizonte − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



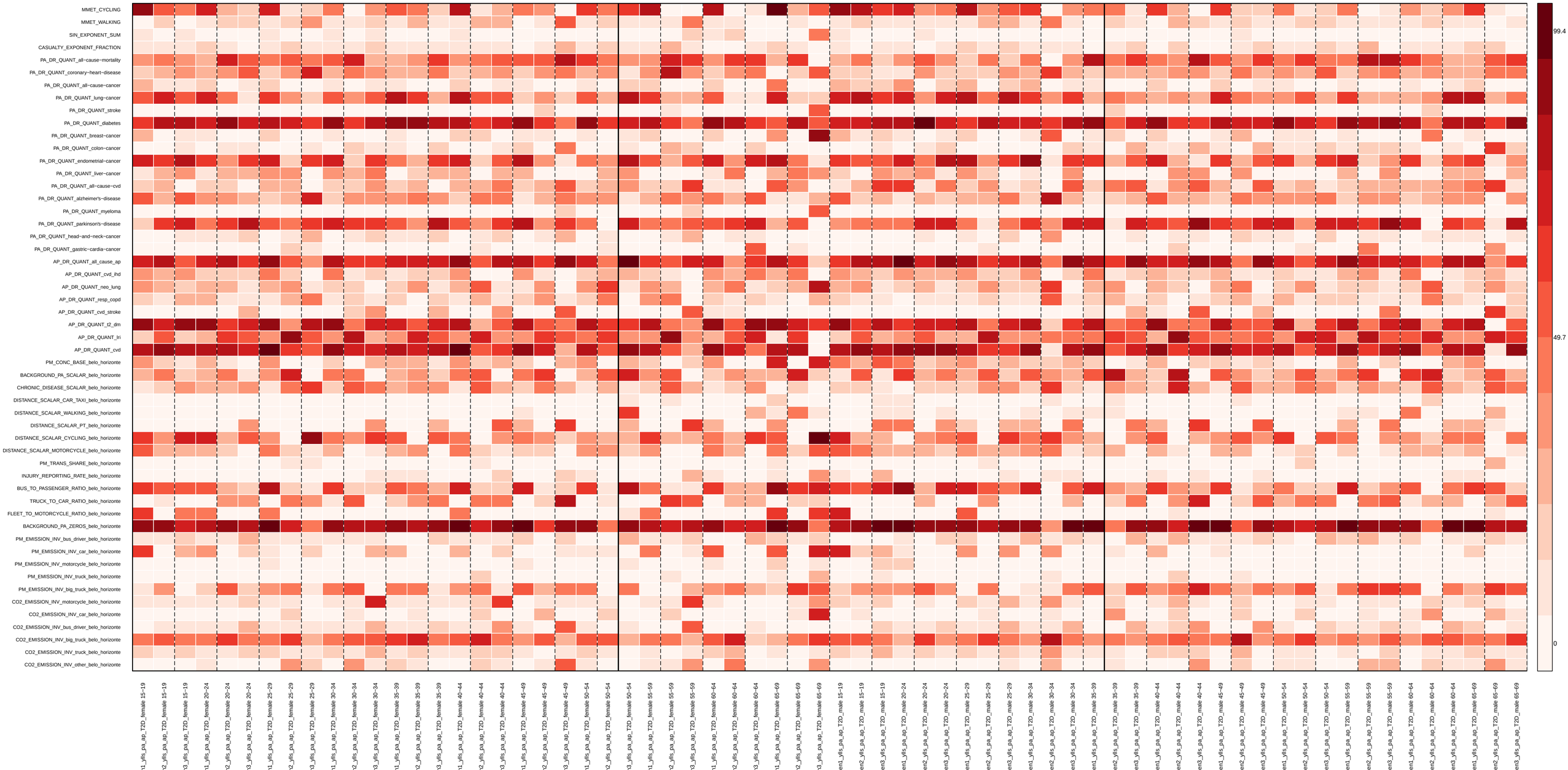
belo_horizonte − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



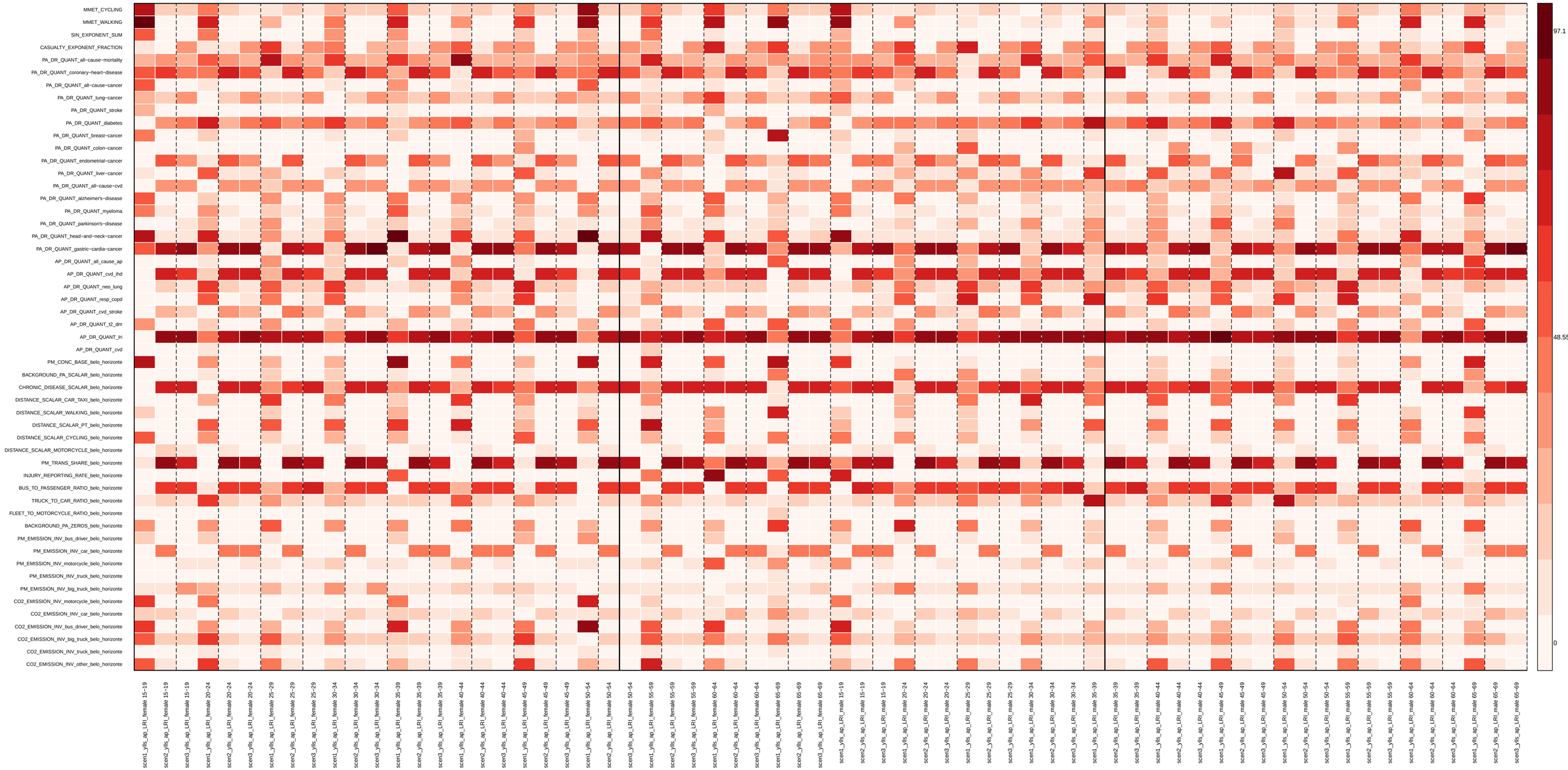
belo_horizonte − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



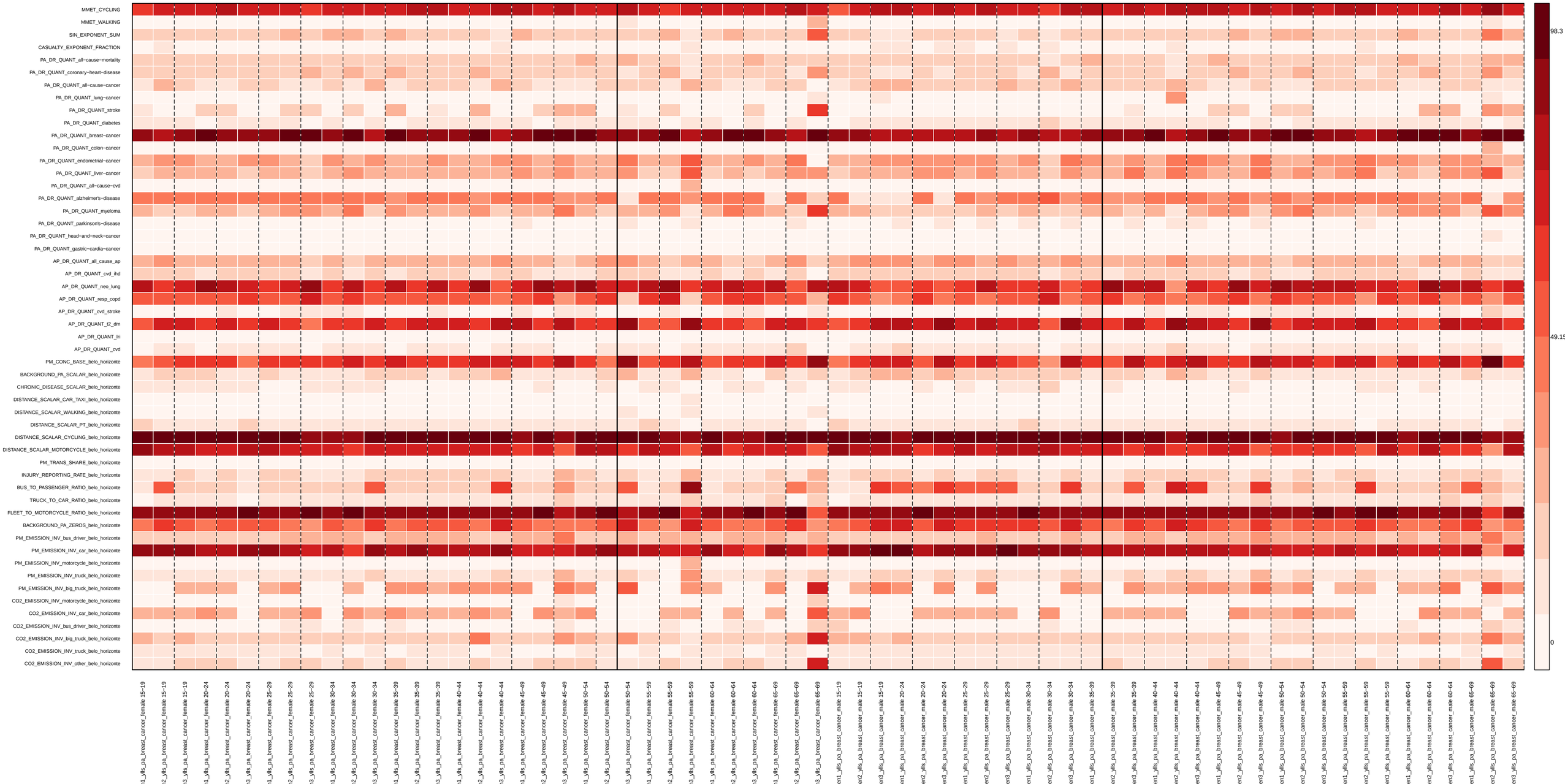
belo_horizonte − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



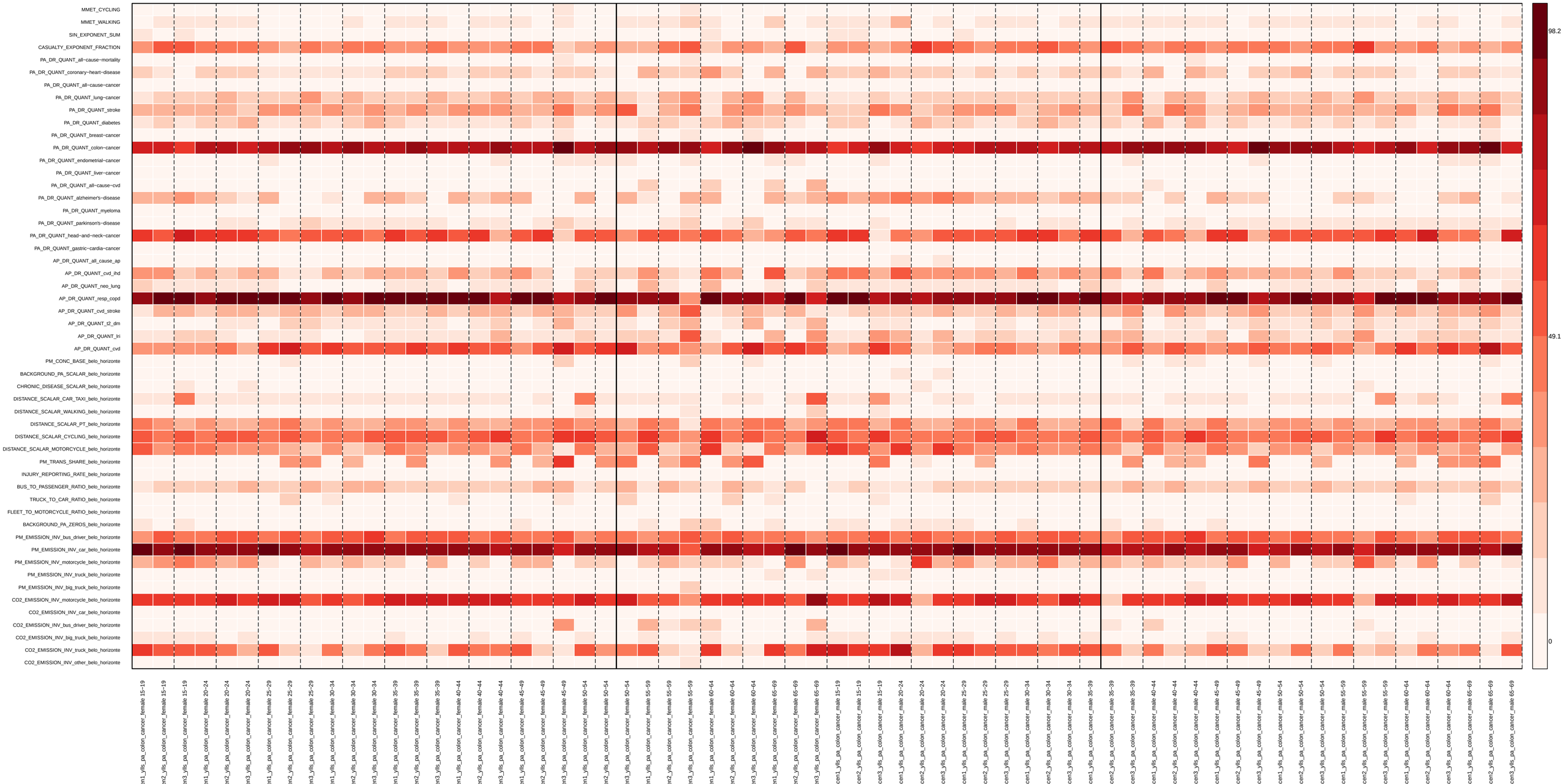
belo_horizonte − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



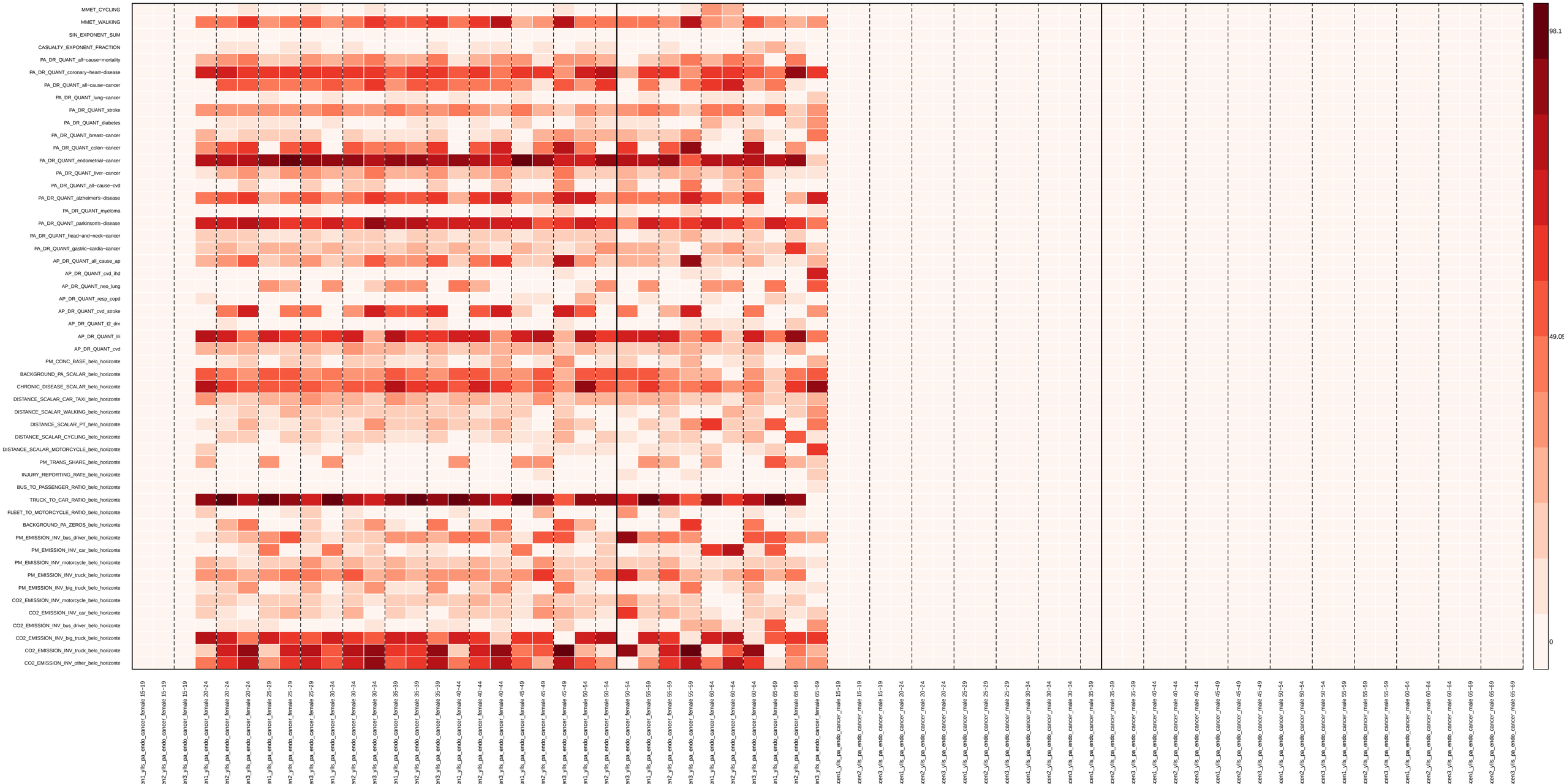
belo_horizonte − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



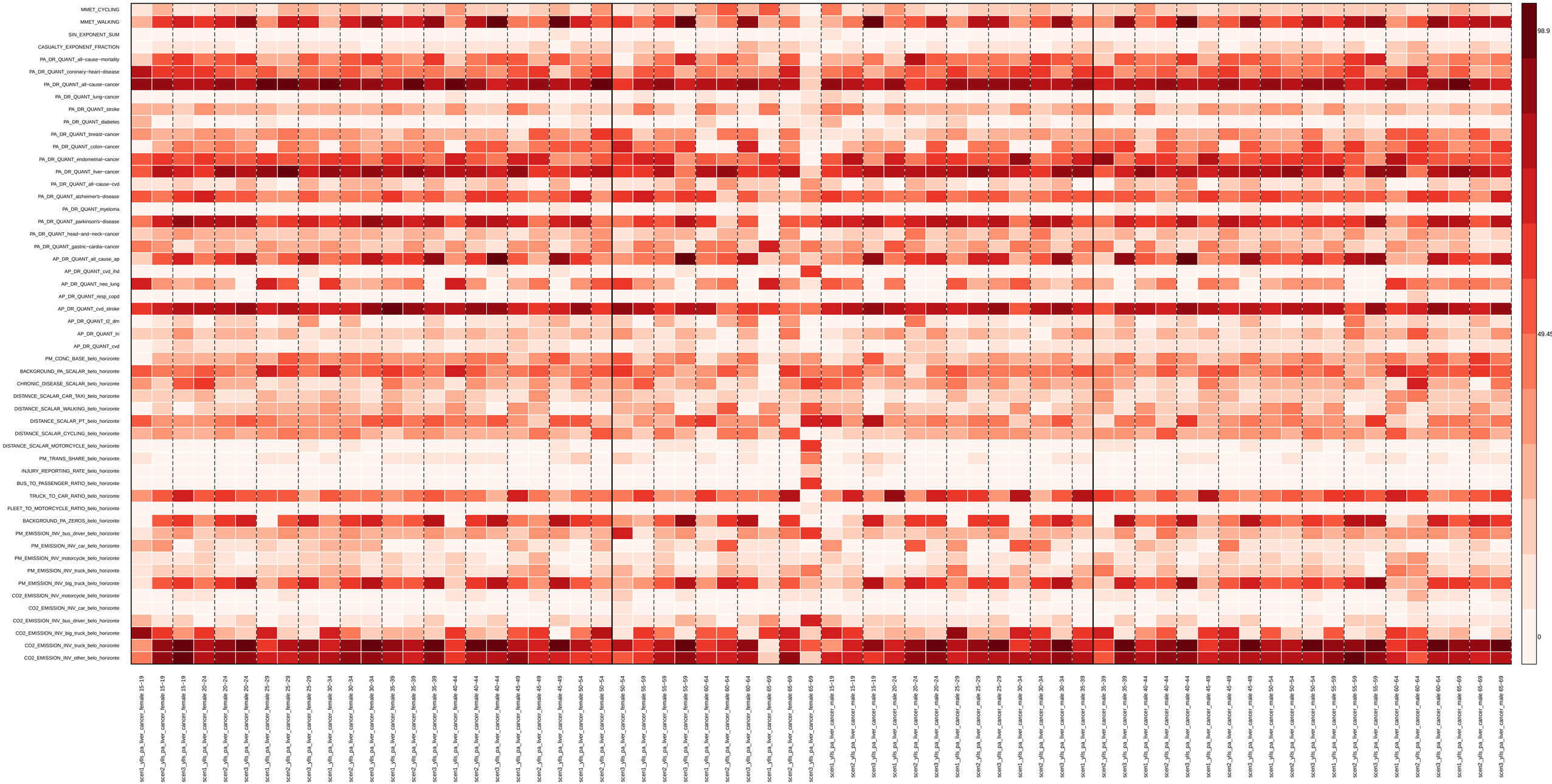
belo_horizonte − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



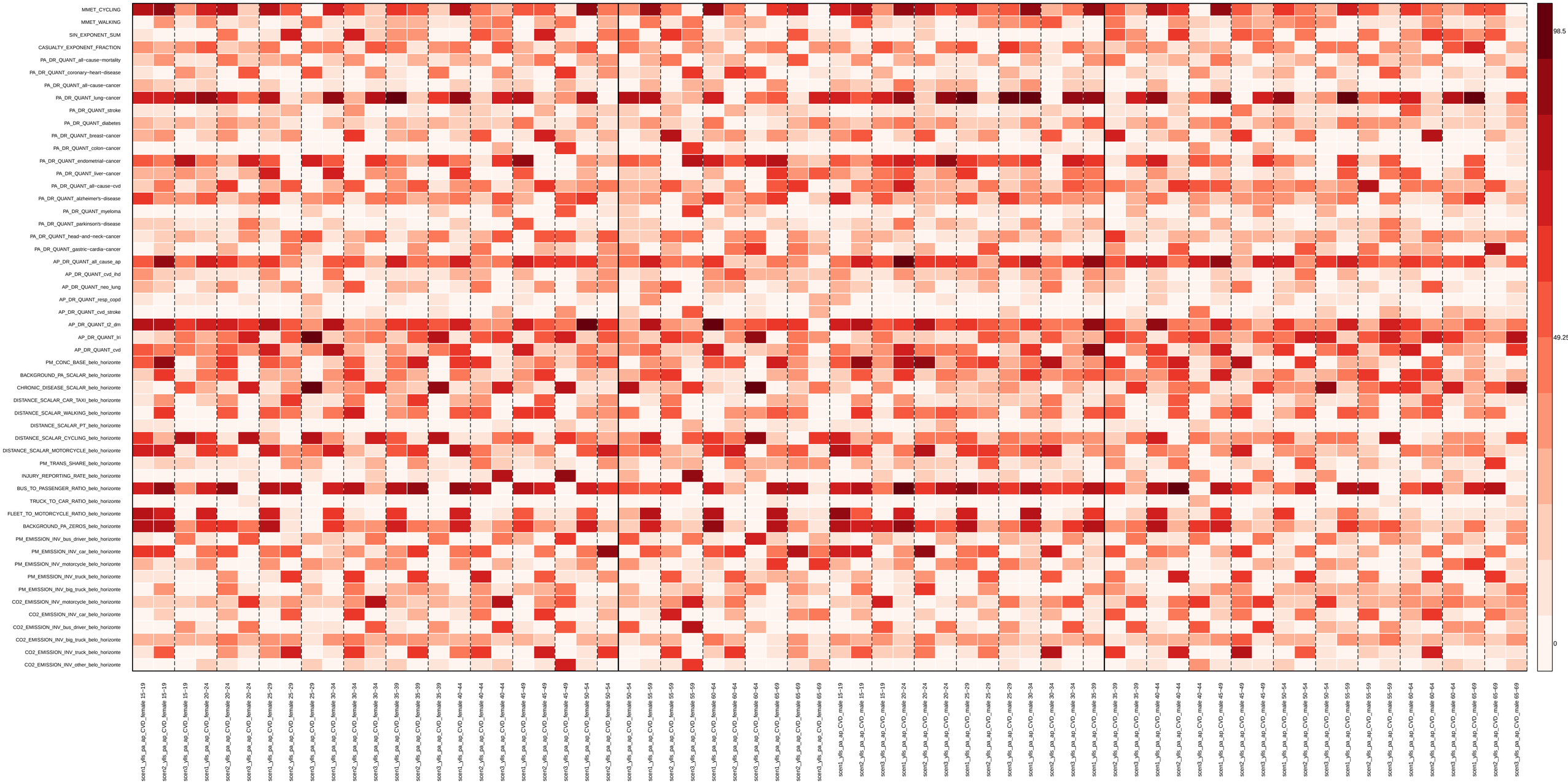
belo_horizonte − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



belo_horizonte − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

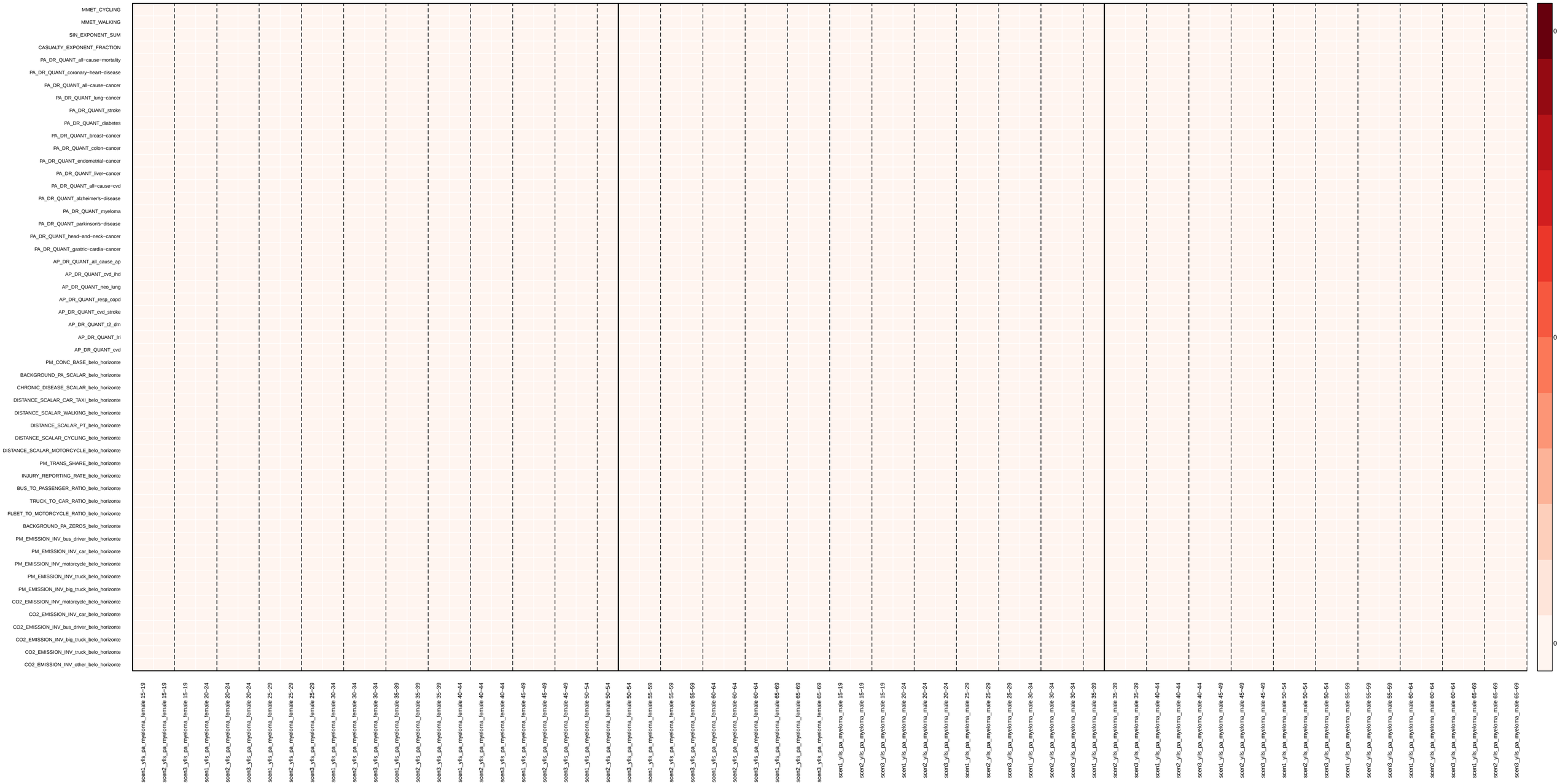


belo_horizonte − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

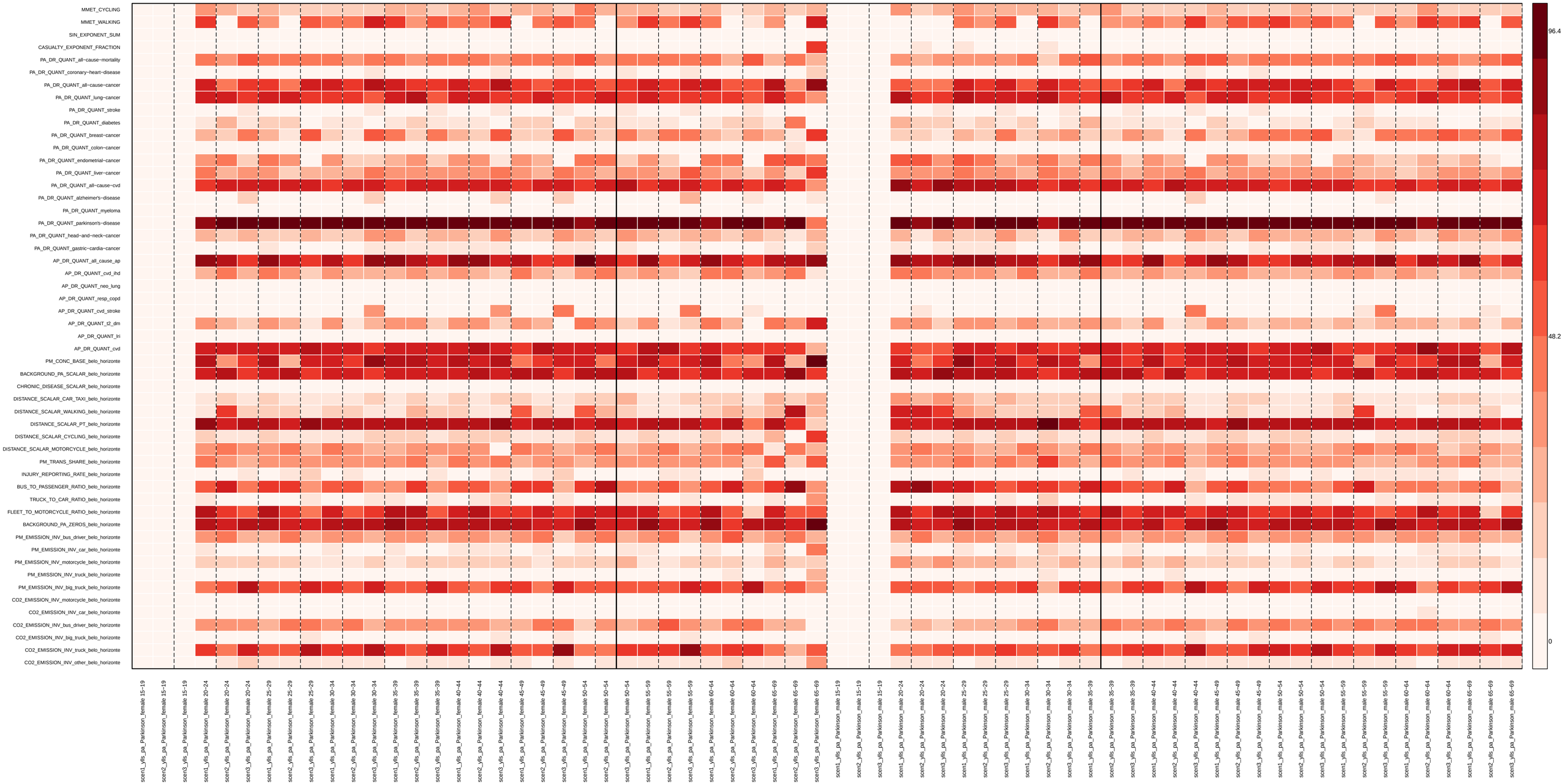


belo_horizonte − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

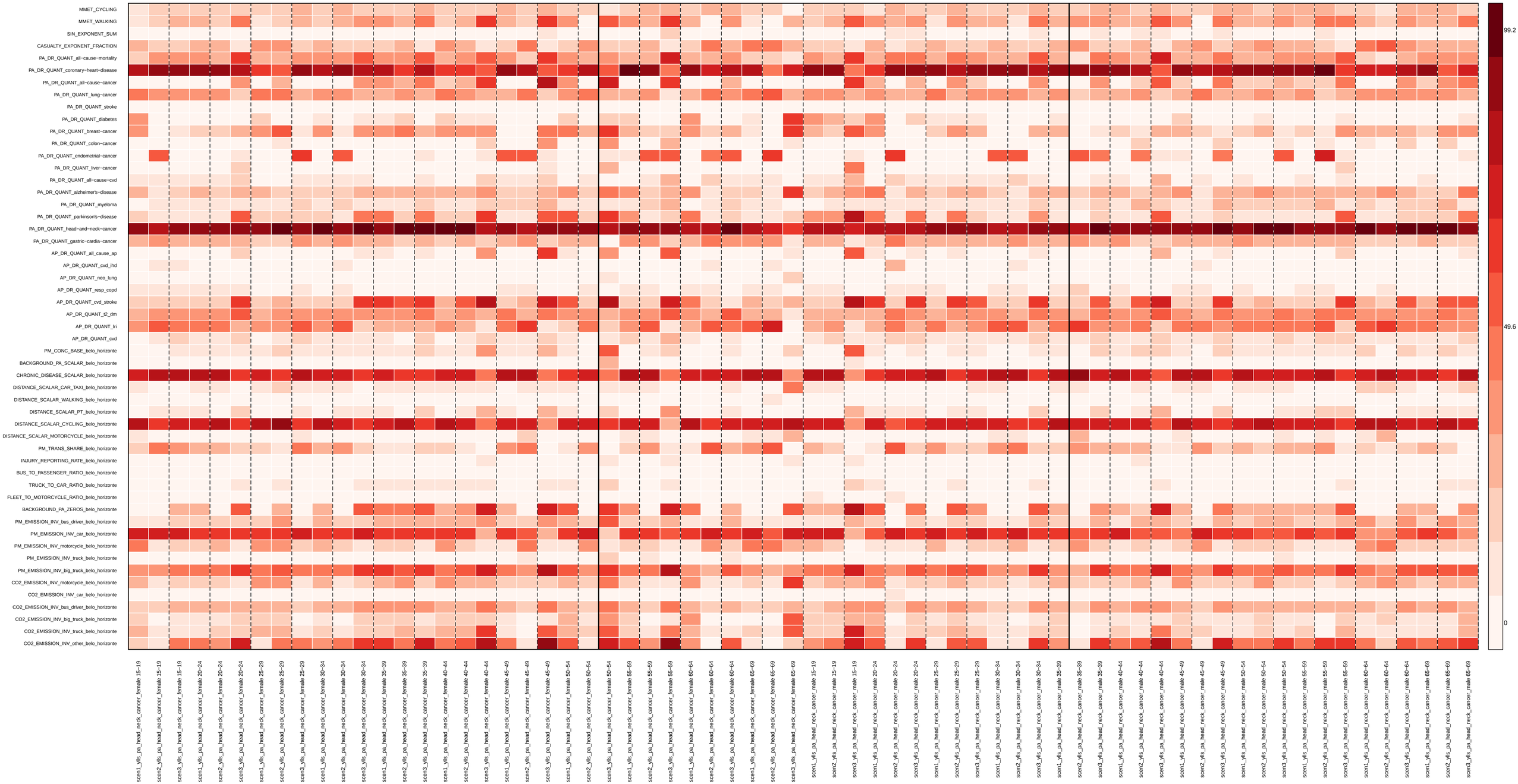
belo_horizonte − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



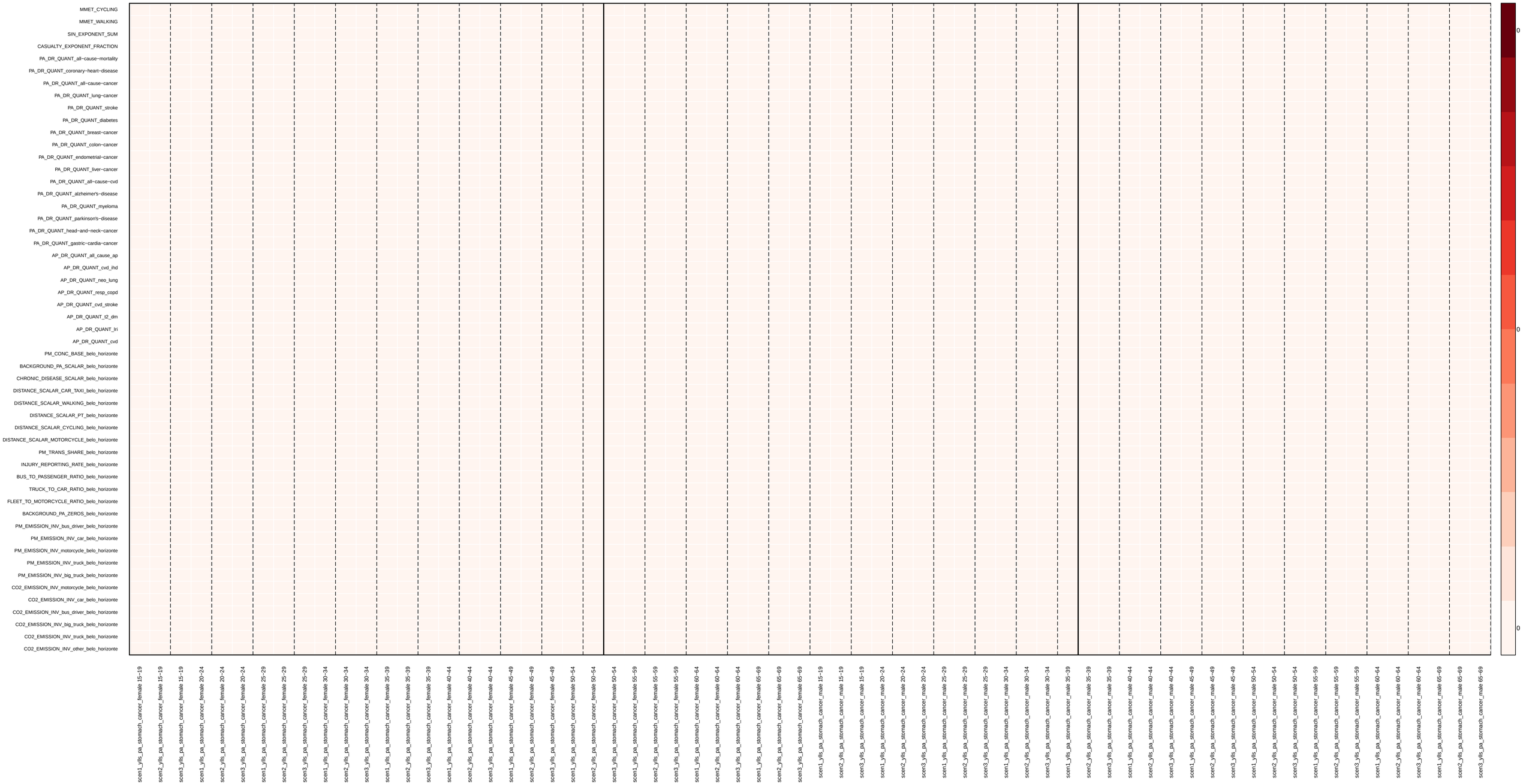
belo_horizonte − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



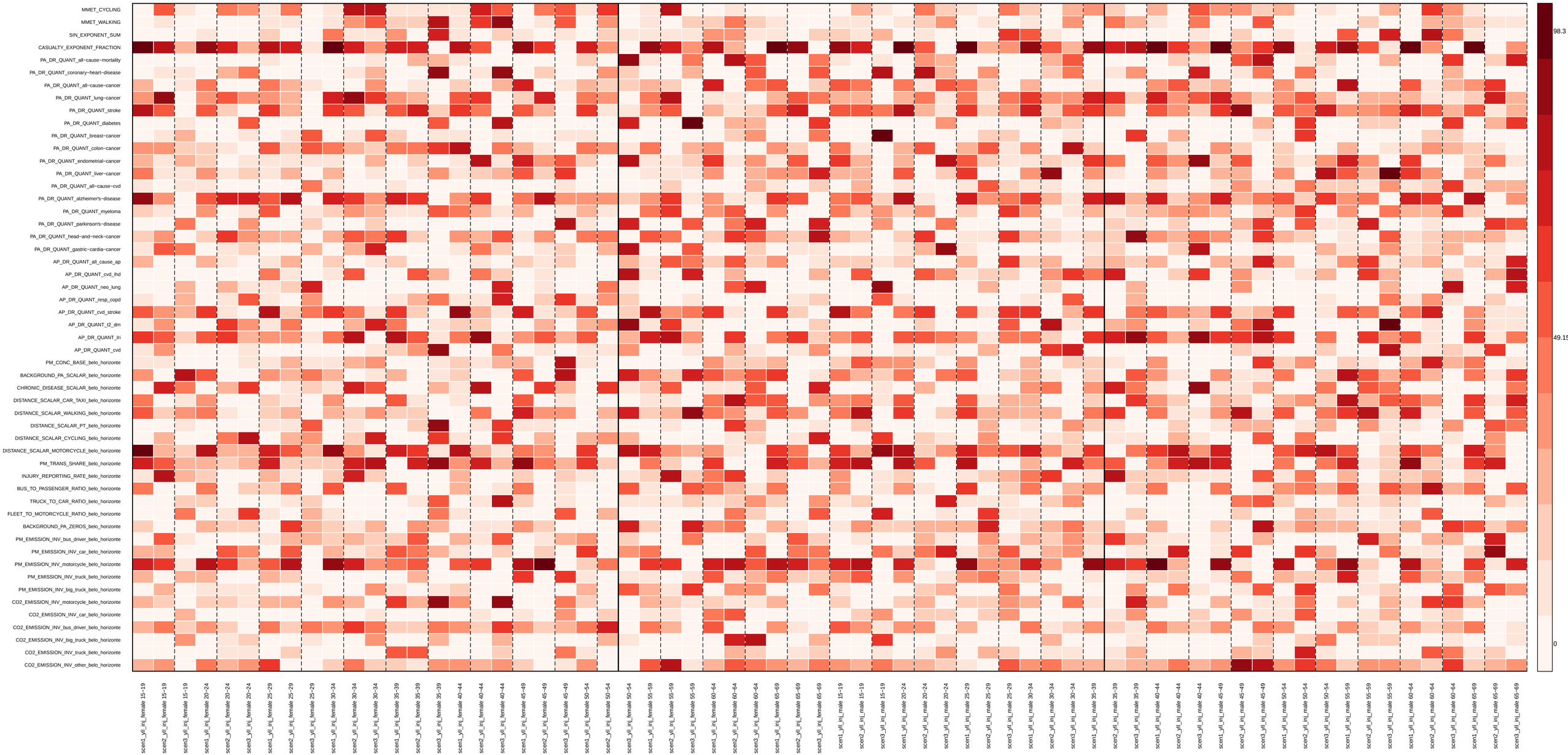
belo_horizonte − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



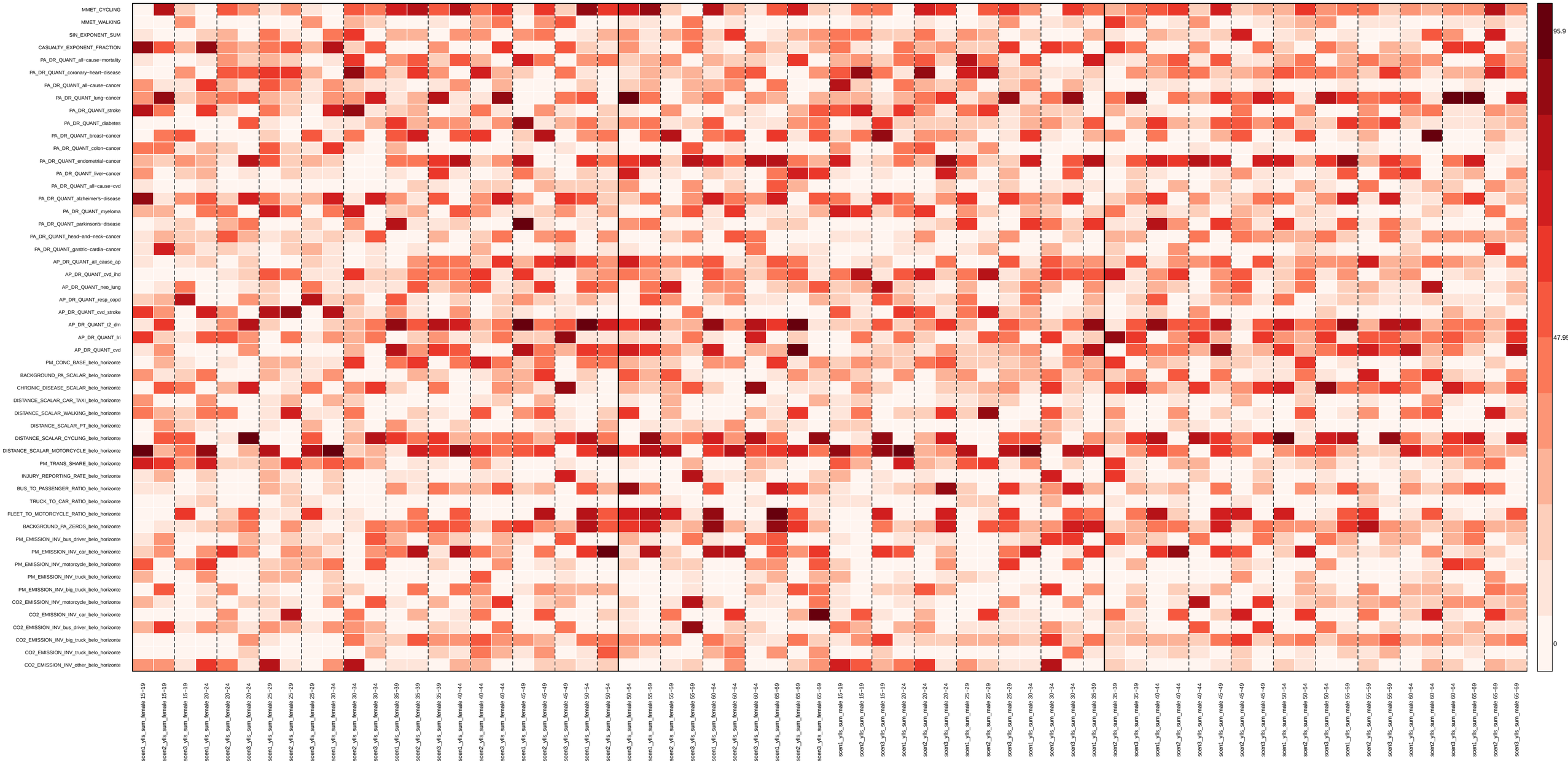
belo_horizonte − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



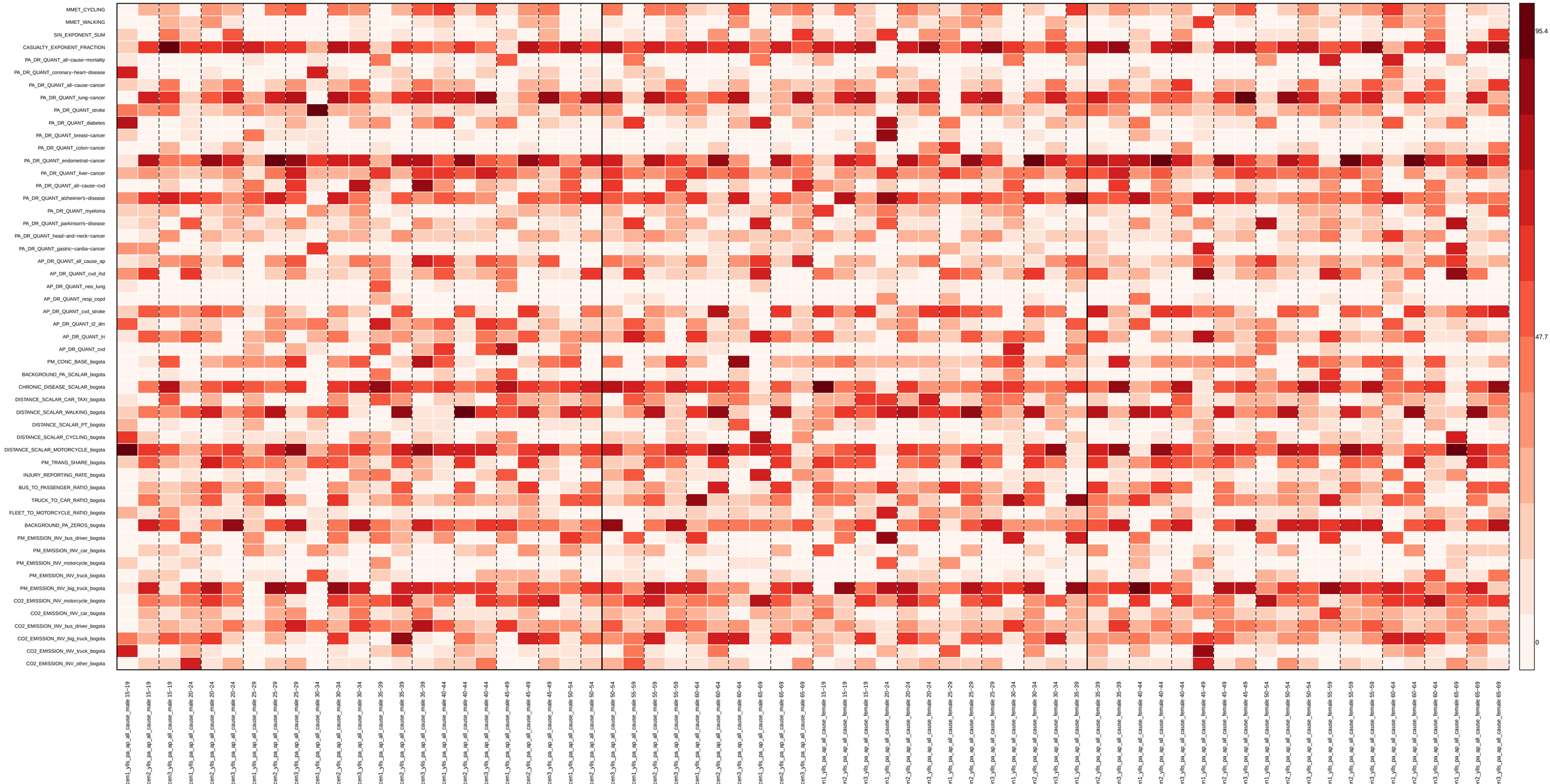
belo_horizonte − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



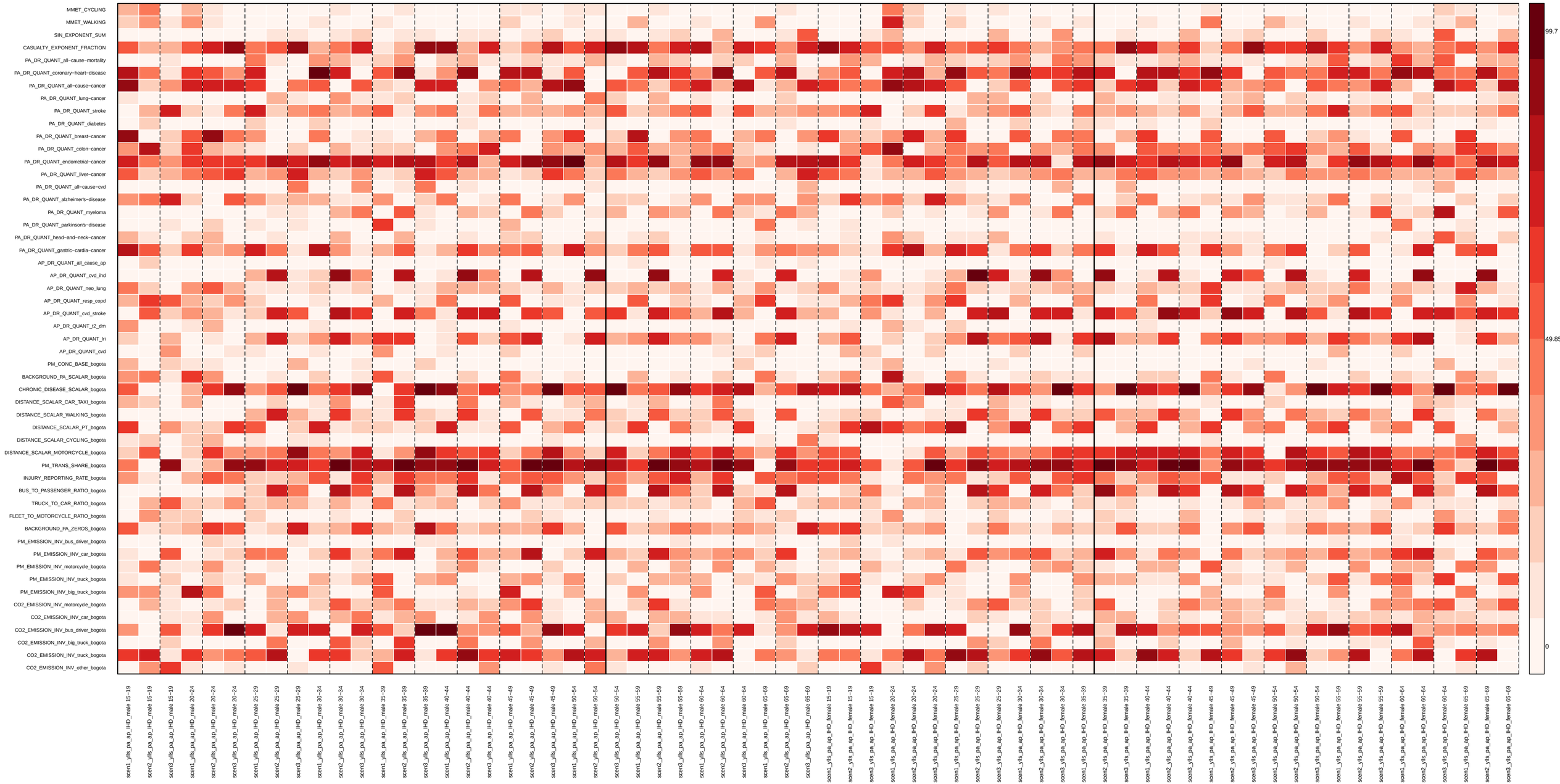
belo_horizonte − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



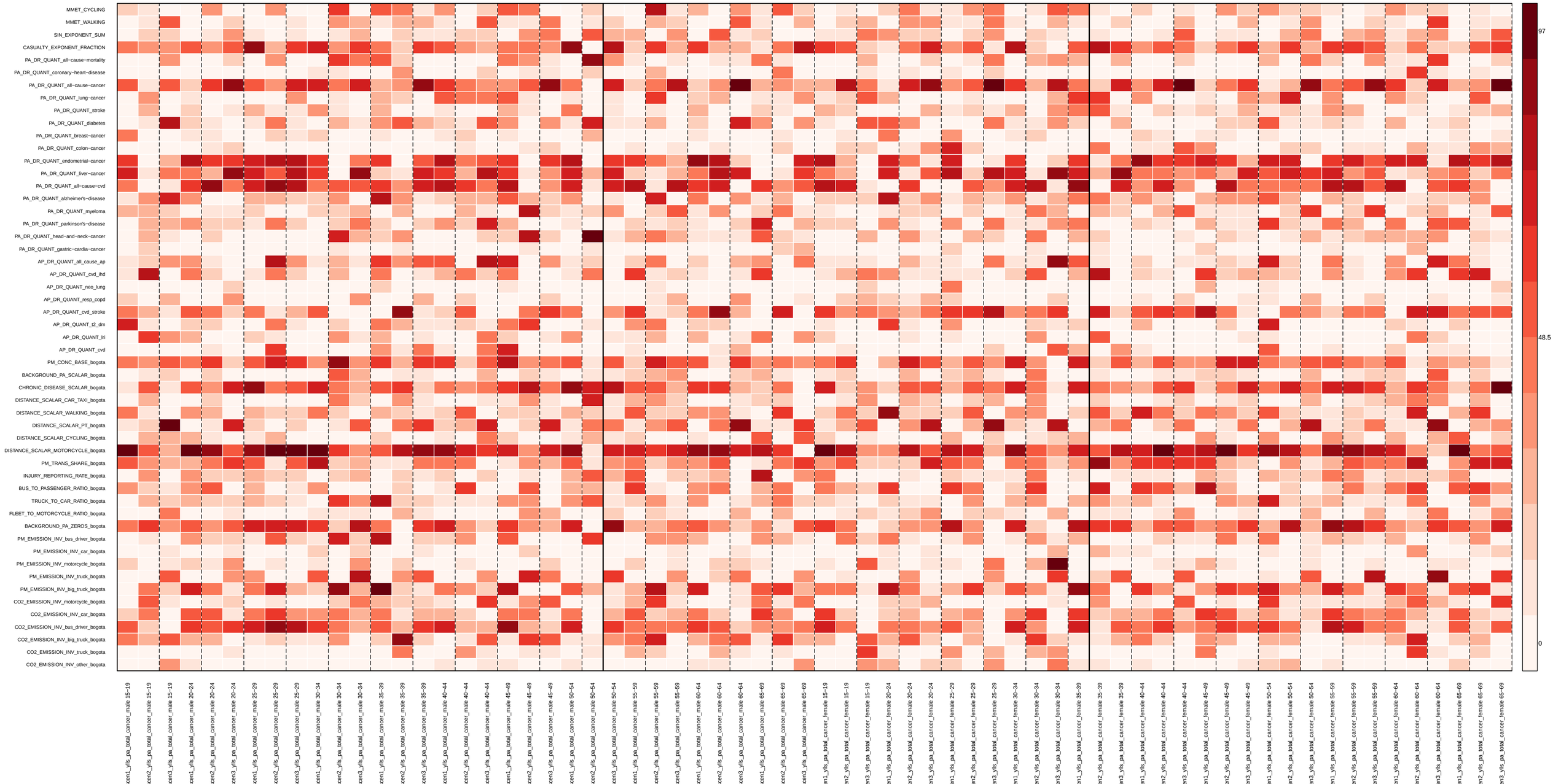
bogota − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



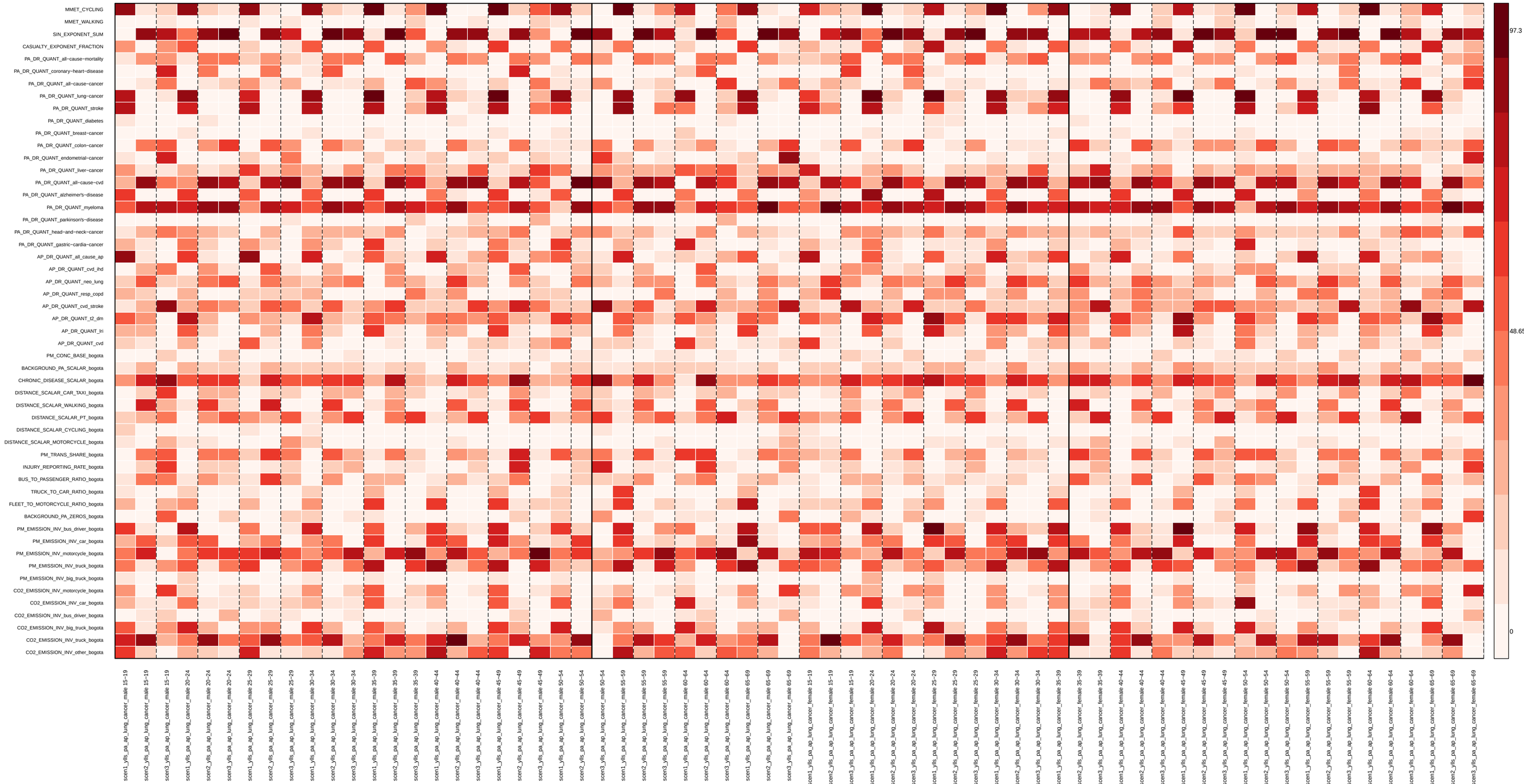
bogota − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



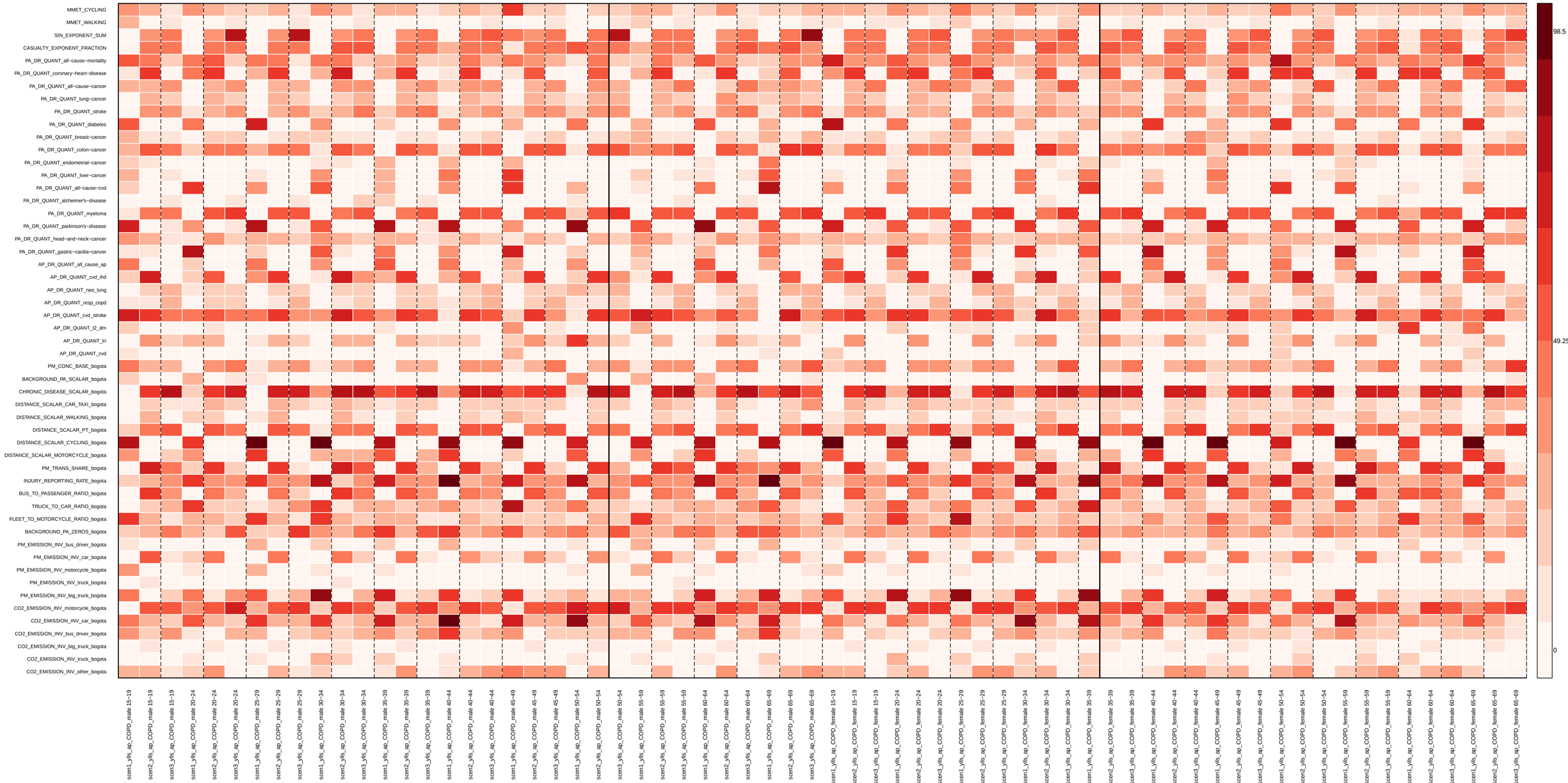
bogota − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



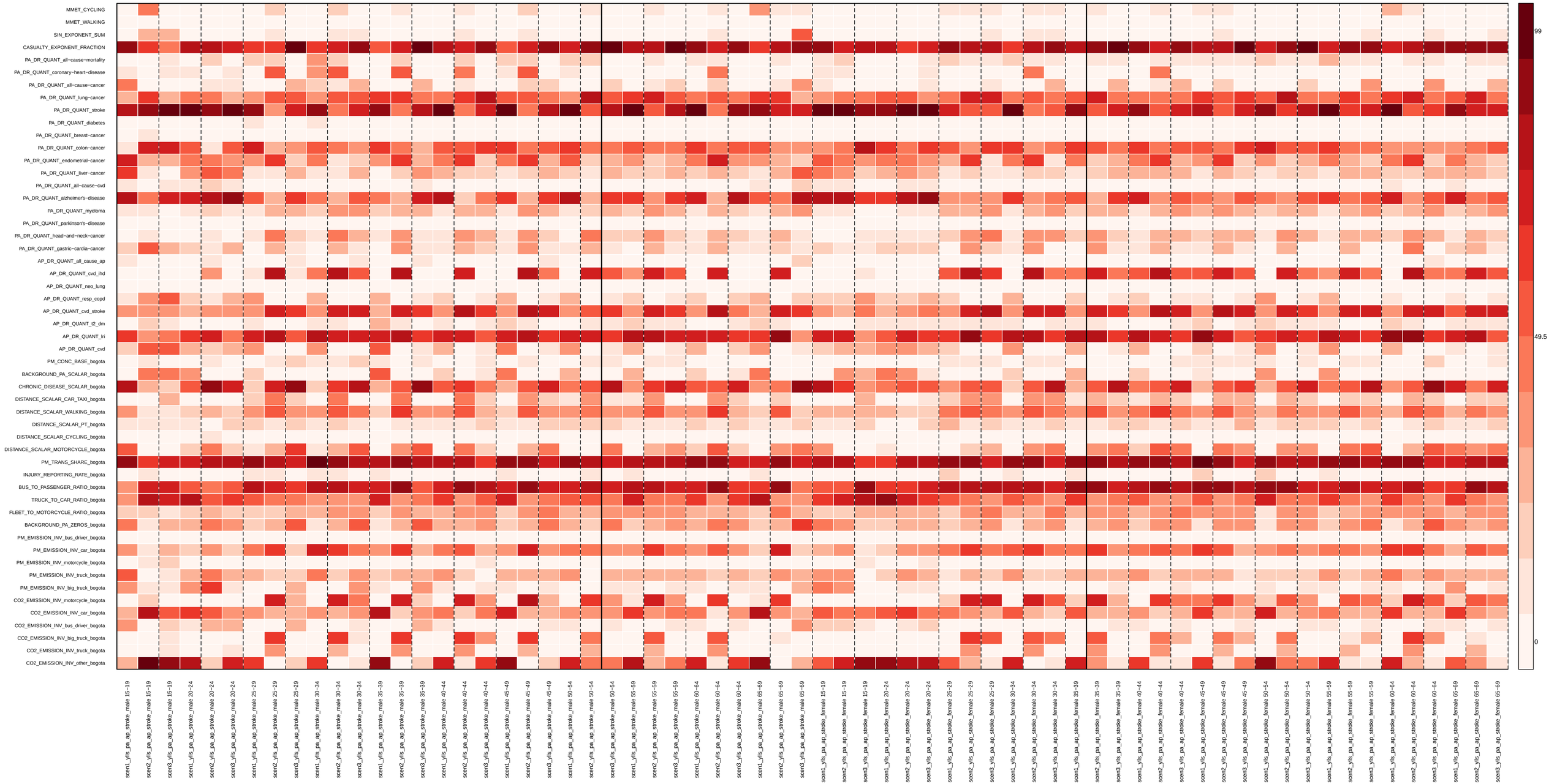
bogota − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



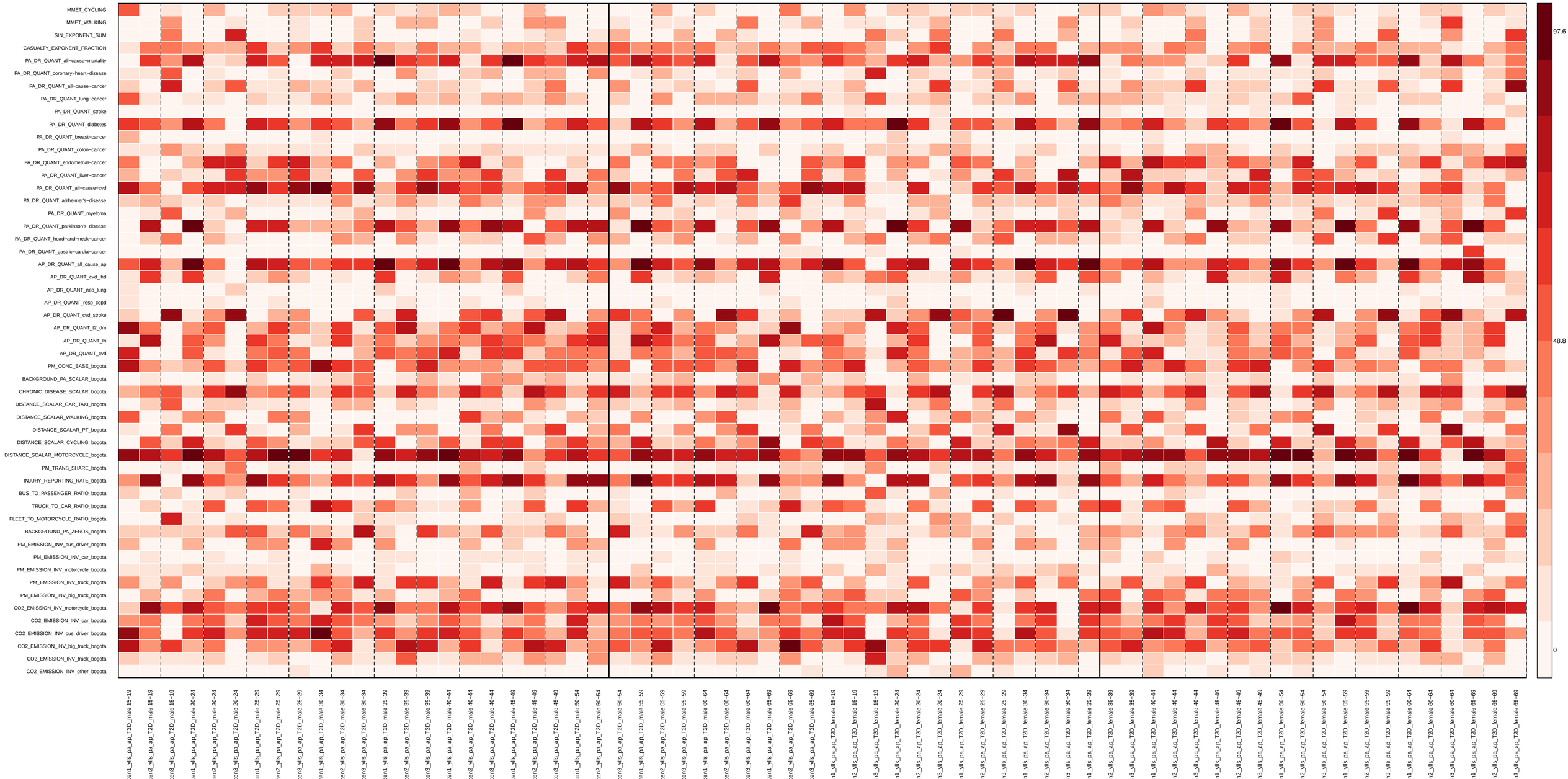
bogota − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



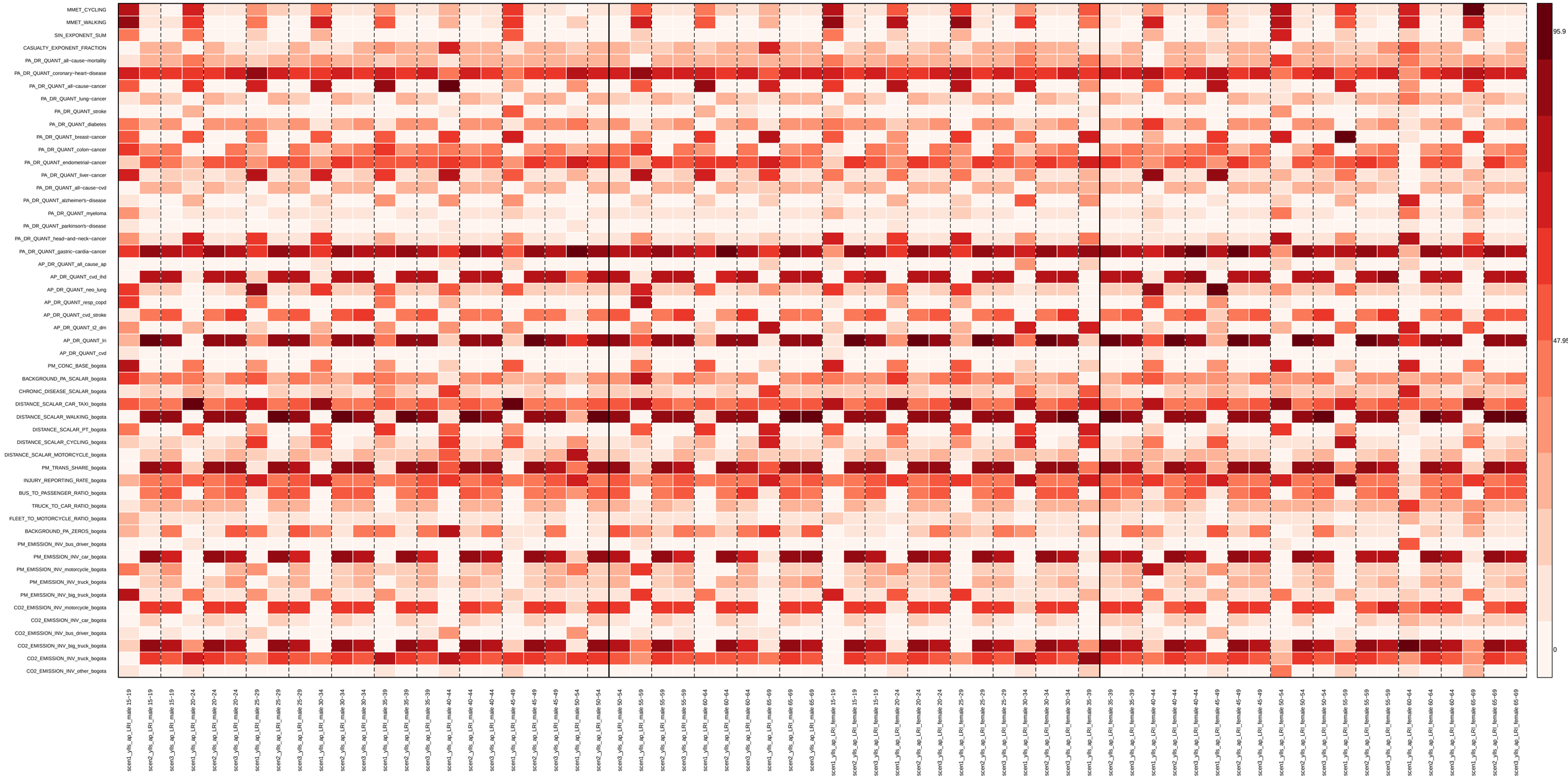
bogota − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



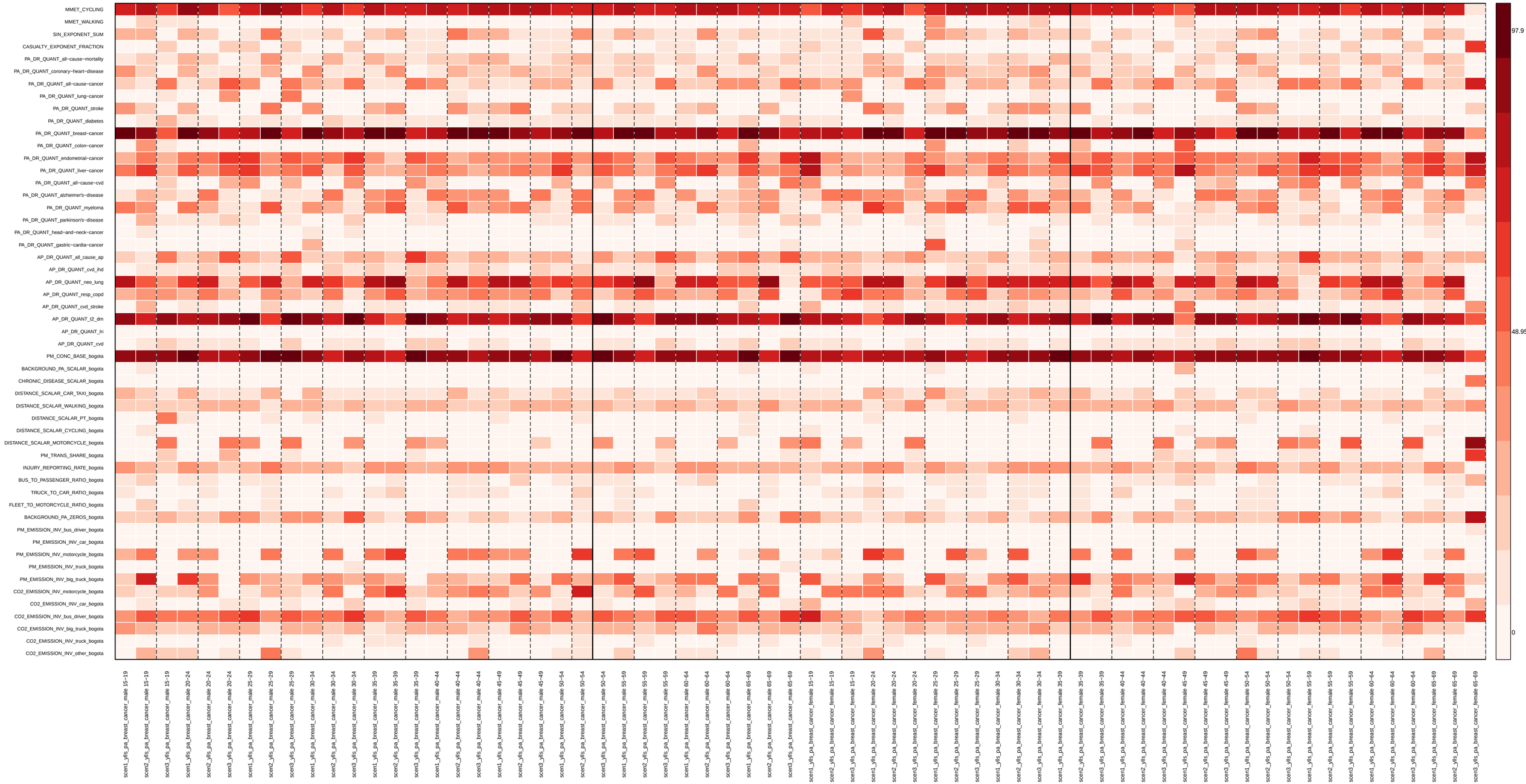
bogota − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



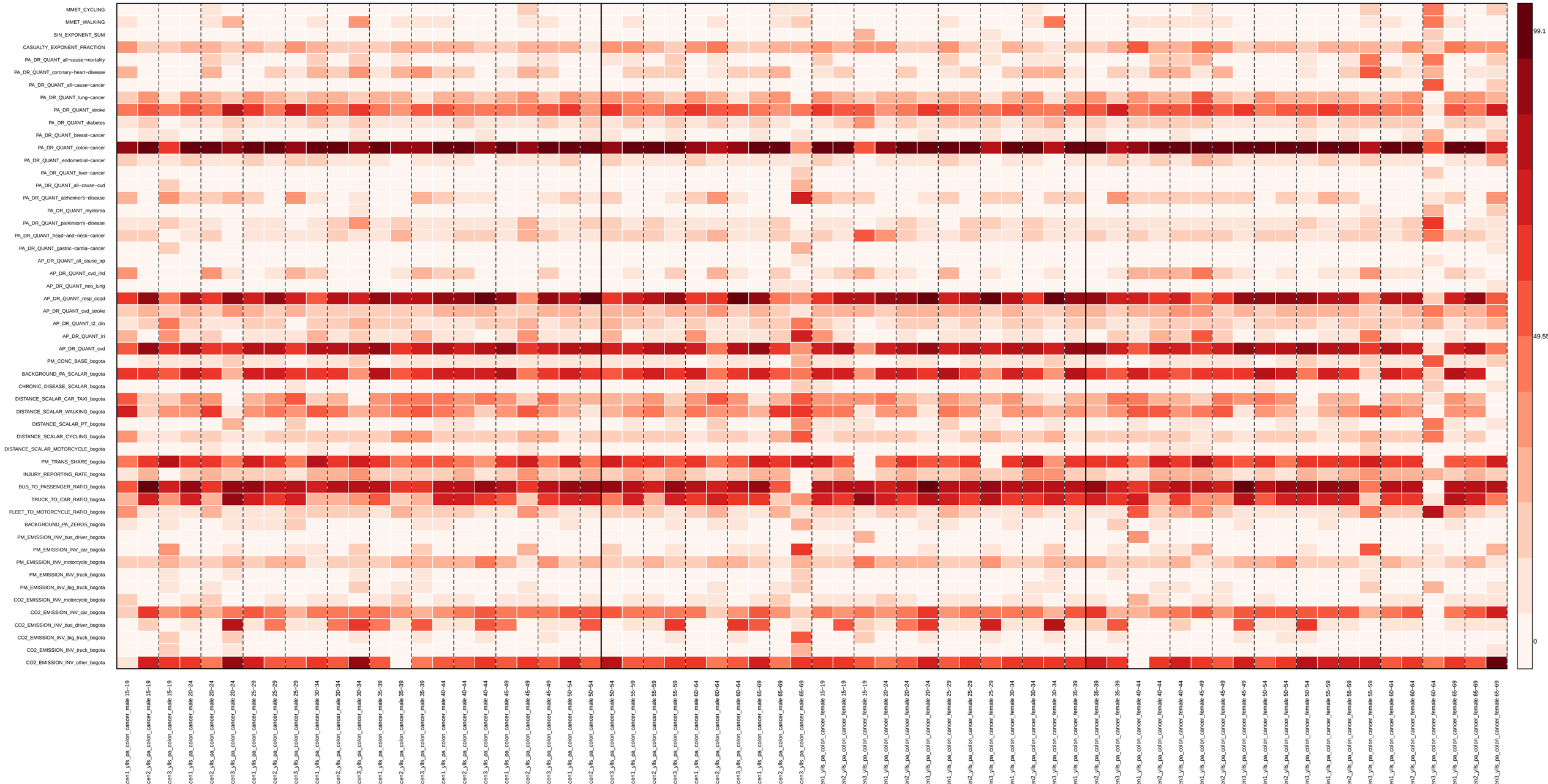
bogota − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



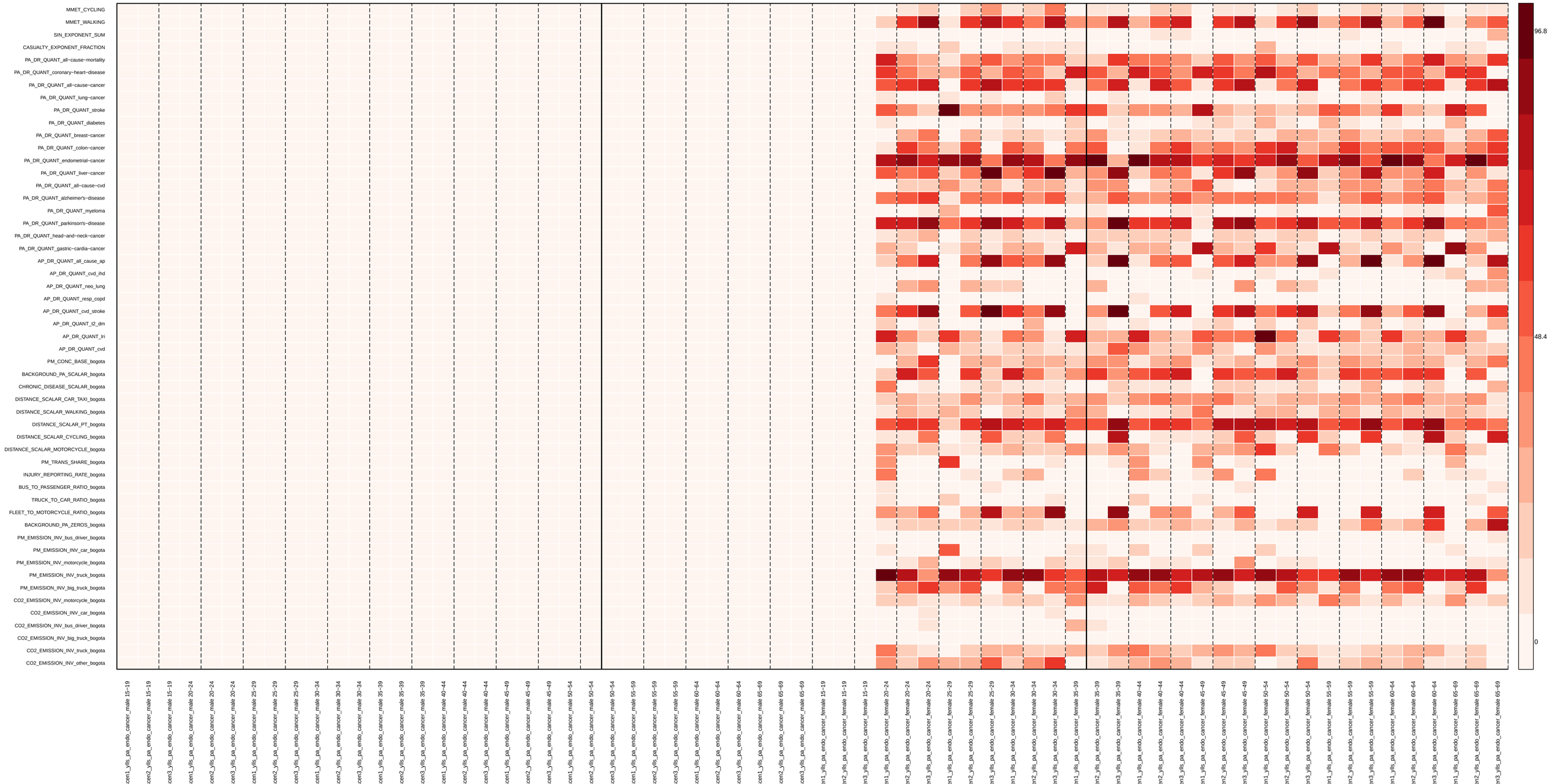
bogota − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



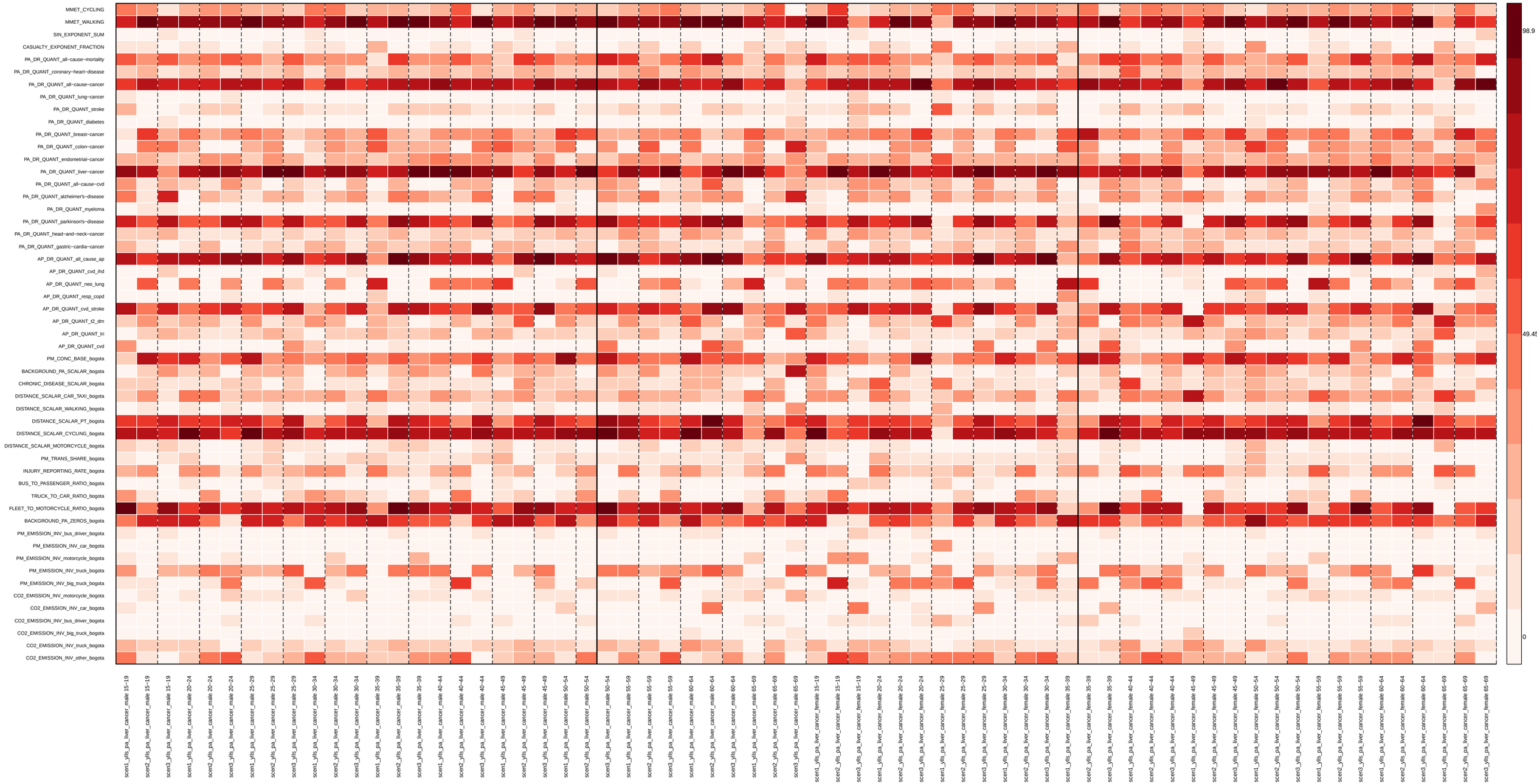
bogota – pa_colon_cancer , No of samples: 10 –
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



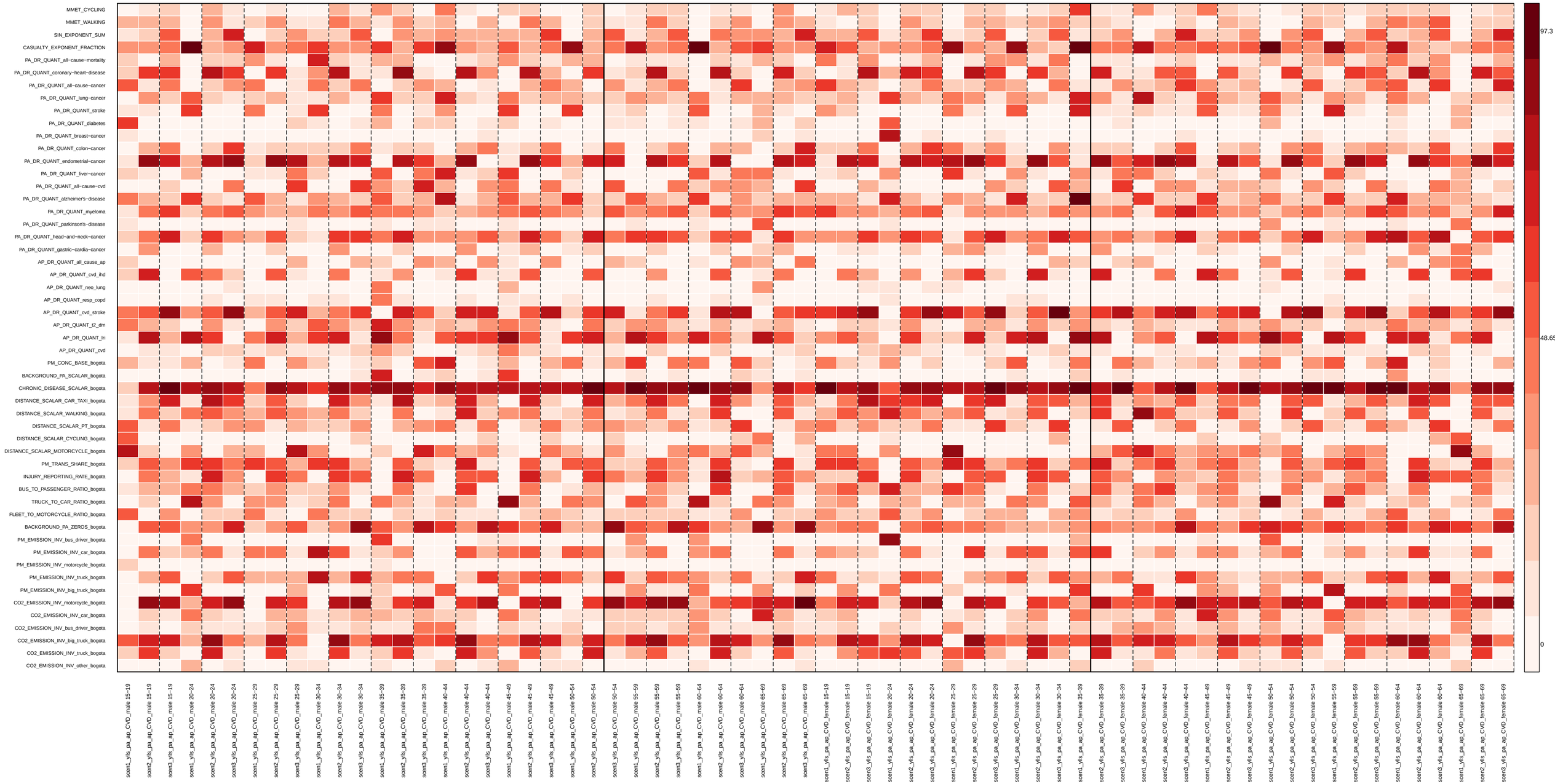
bogota − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



bogota − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

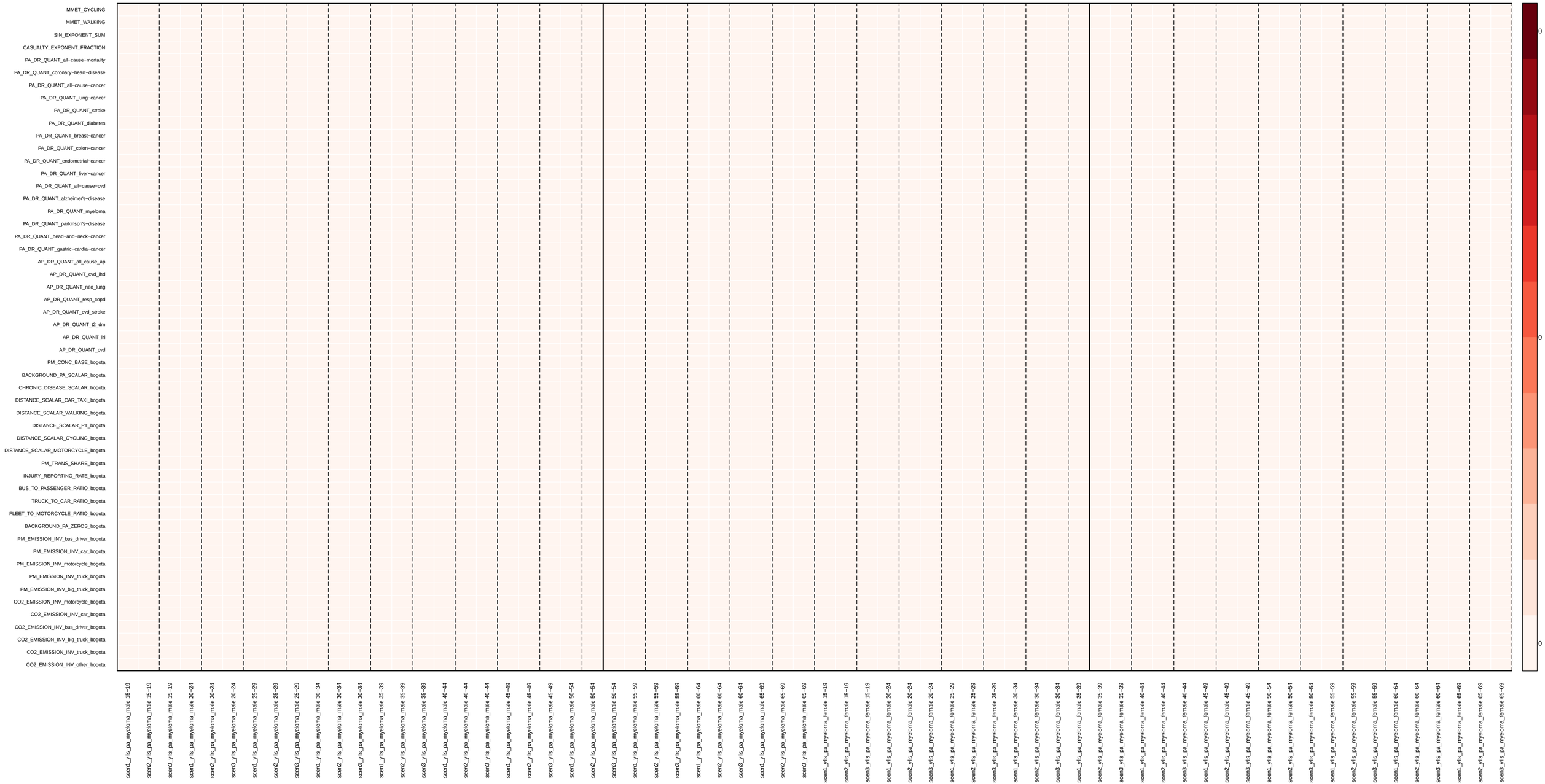


bogota − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

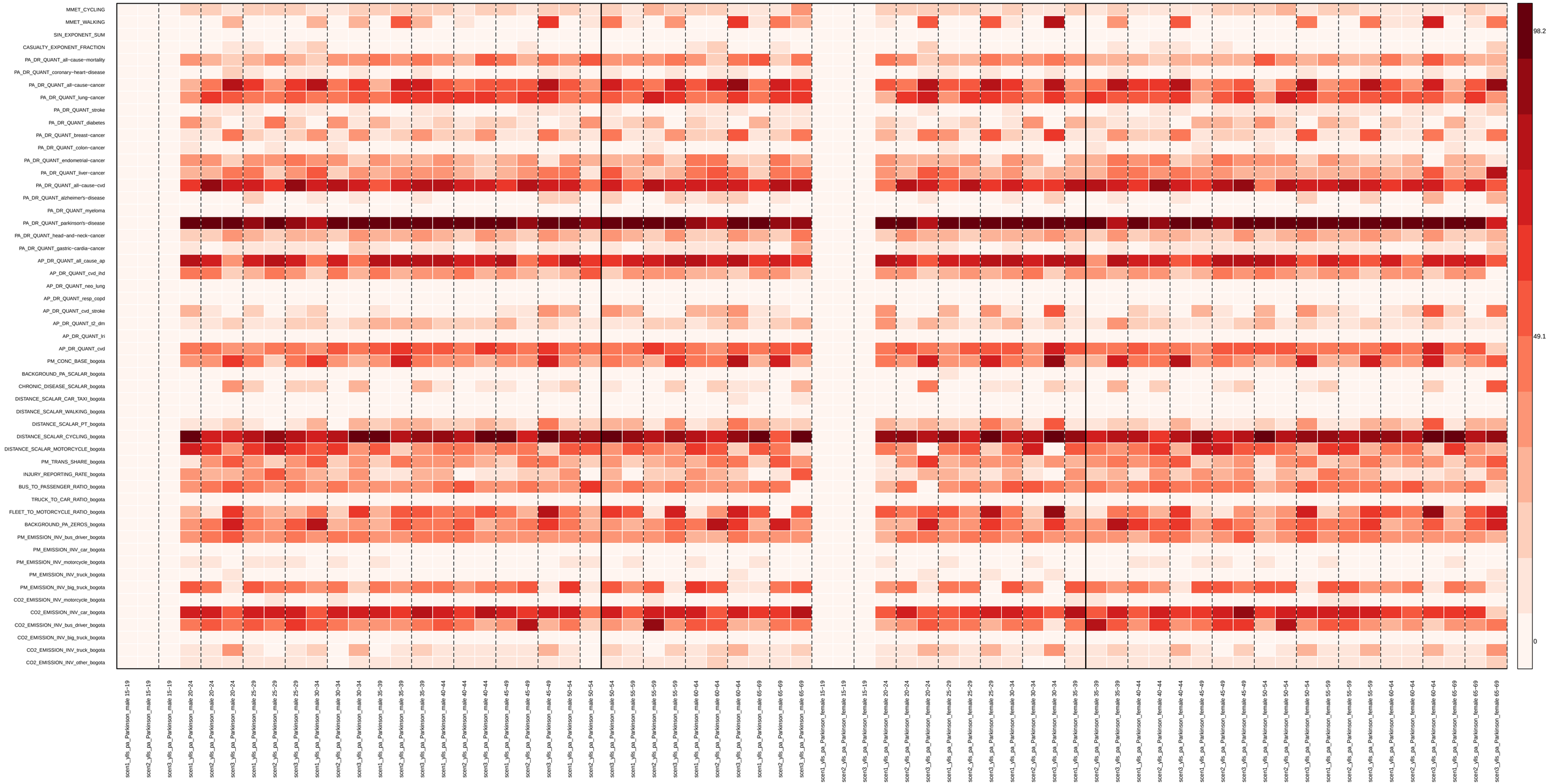


bogota − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

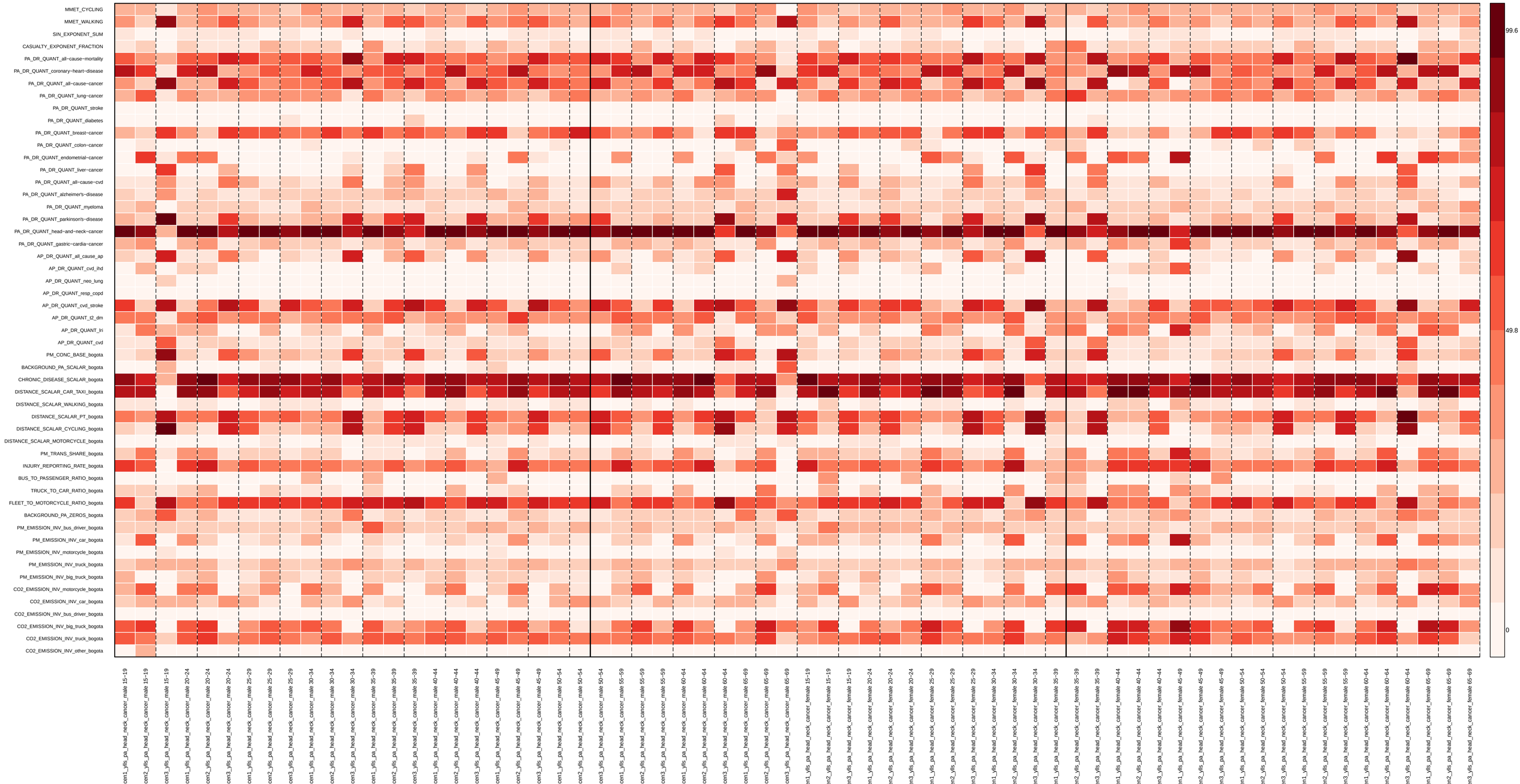
bogota − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



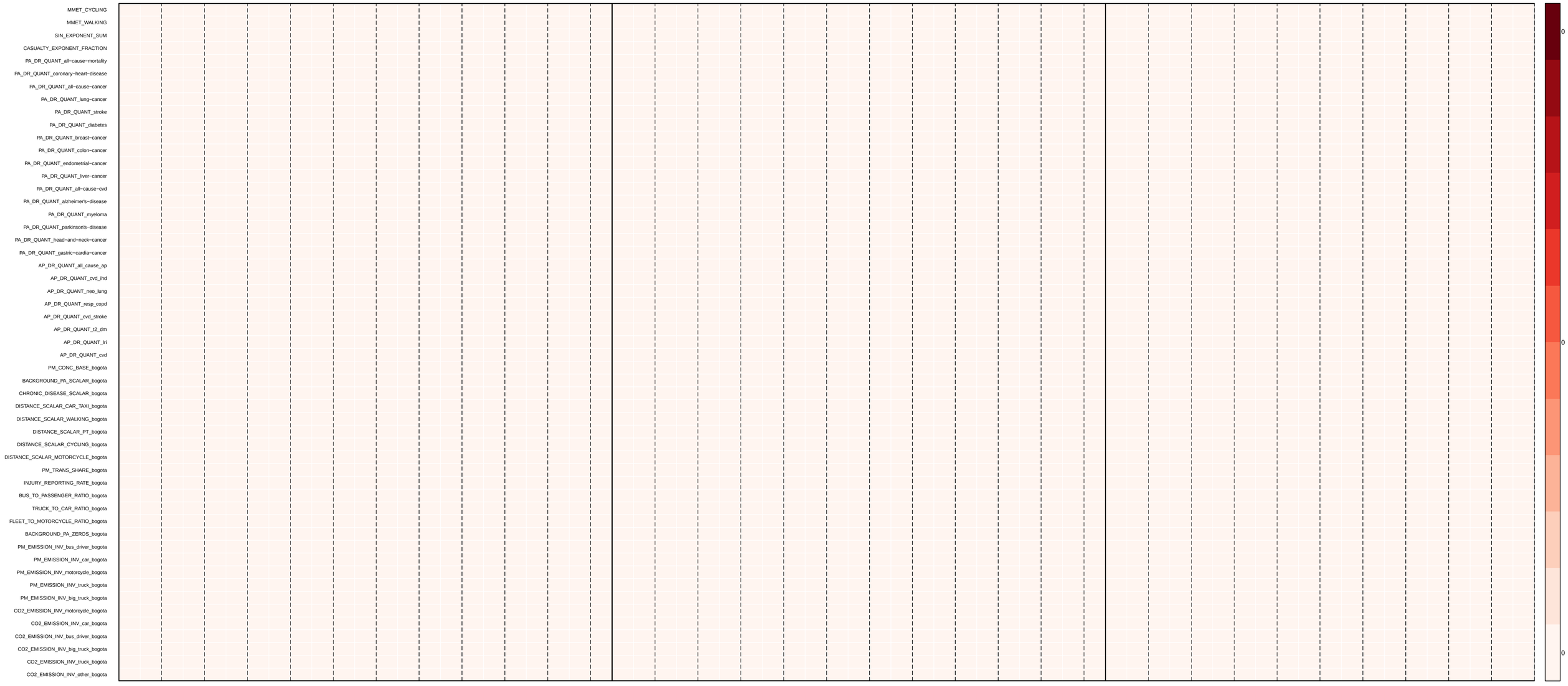
bogota − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



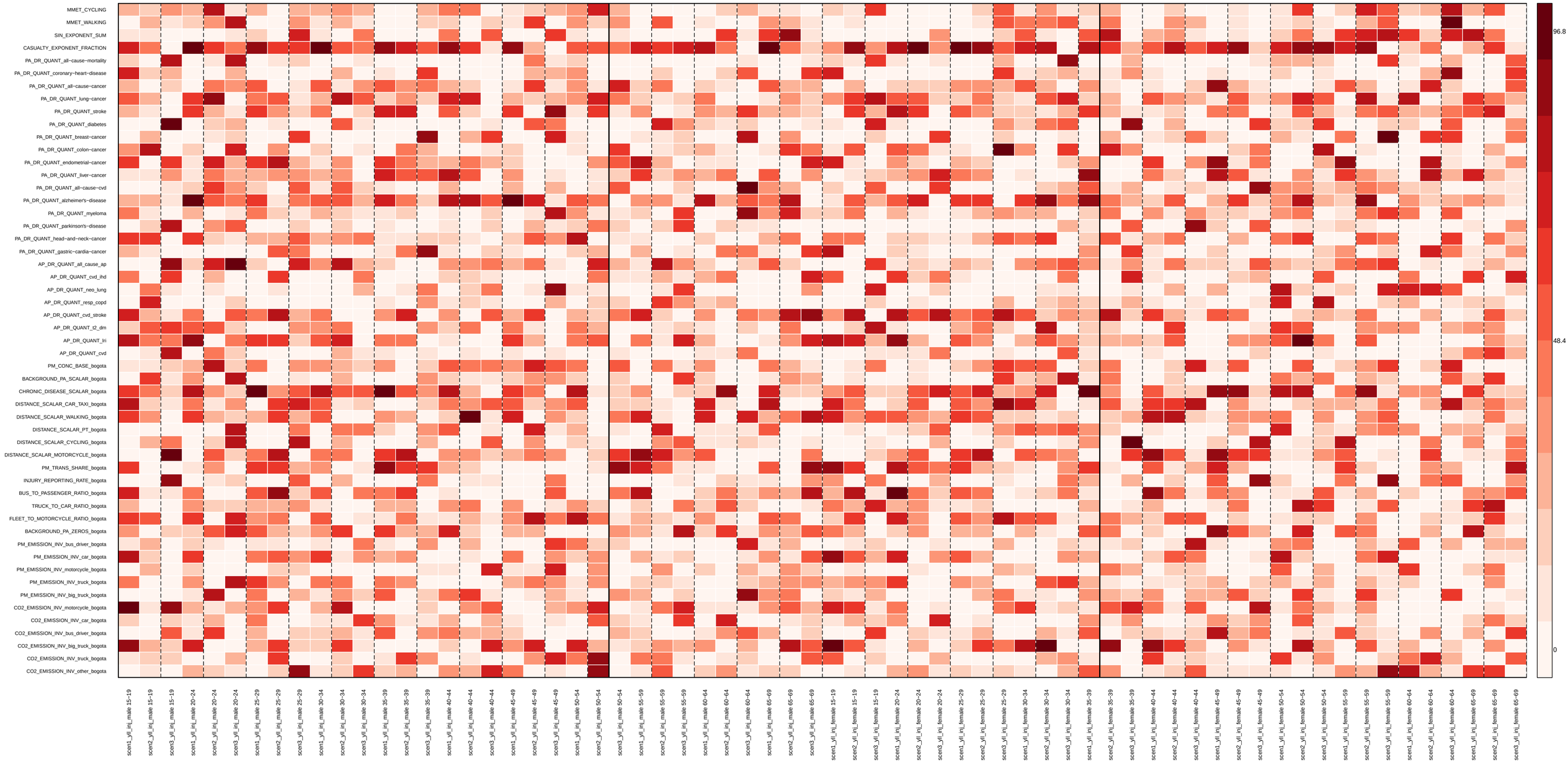
bogota − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



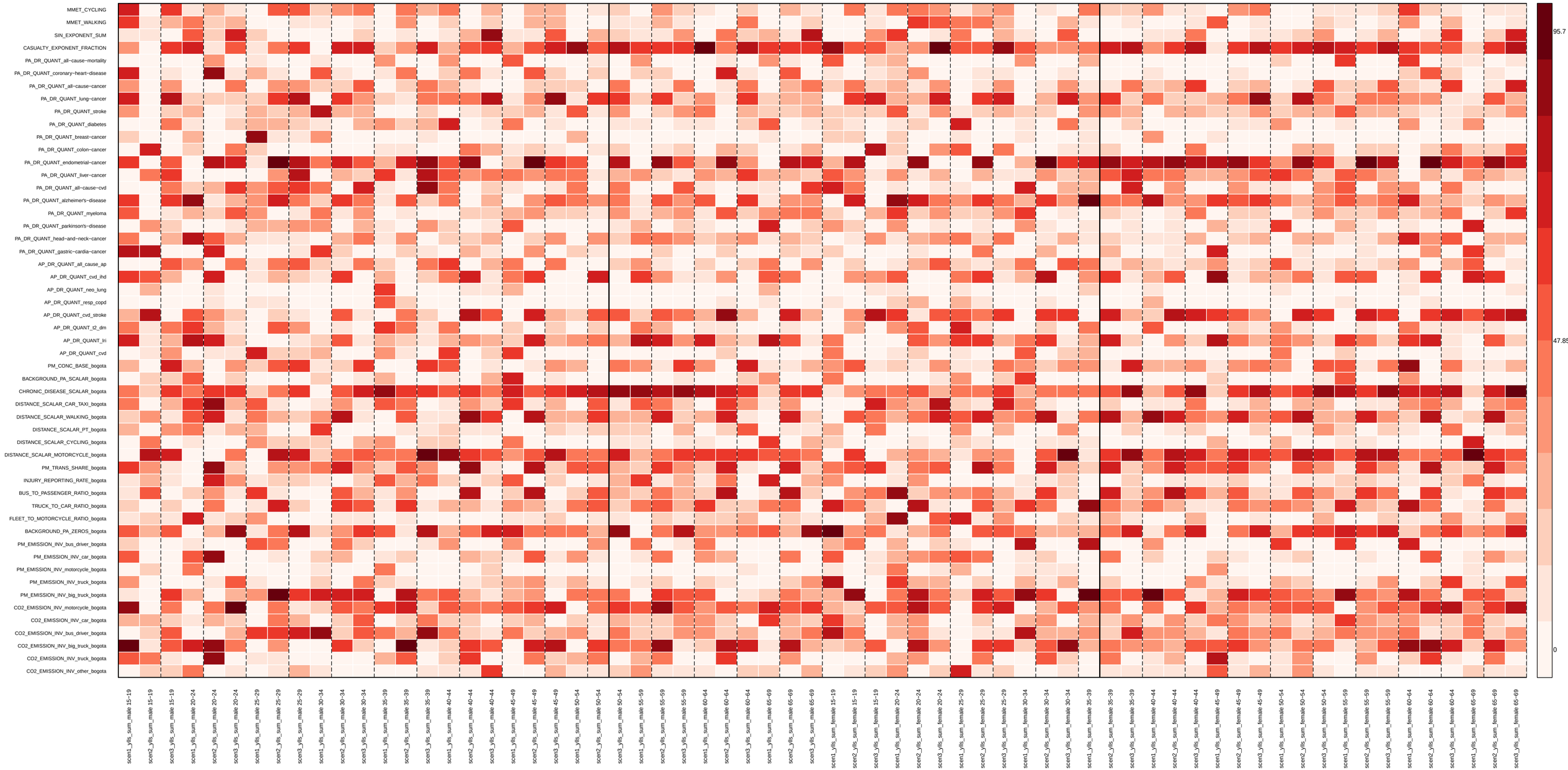
bogota − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



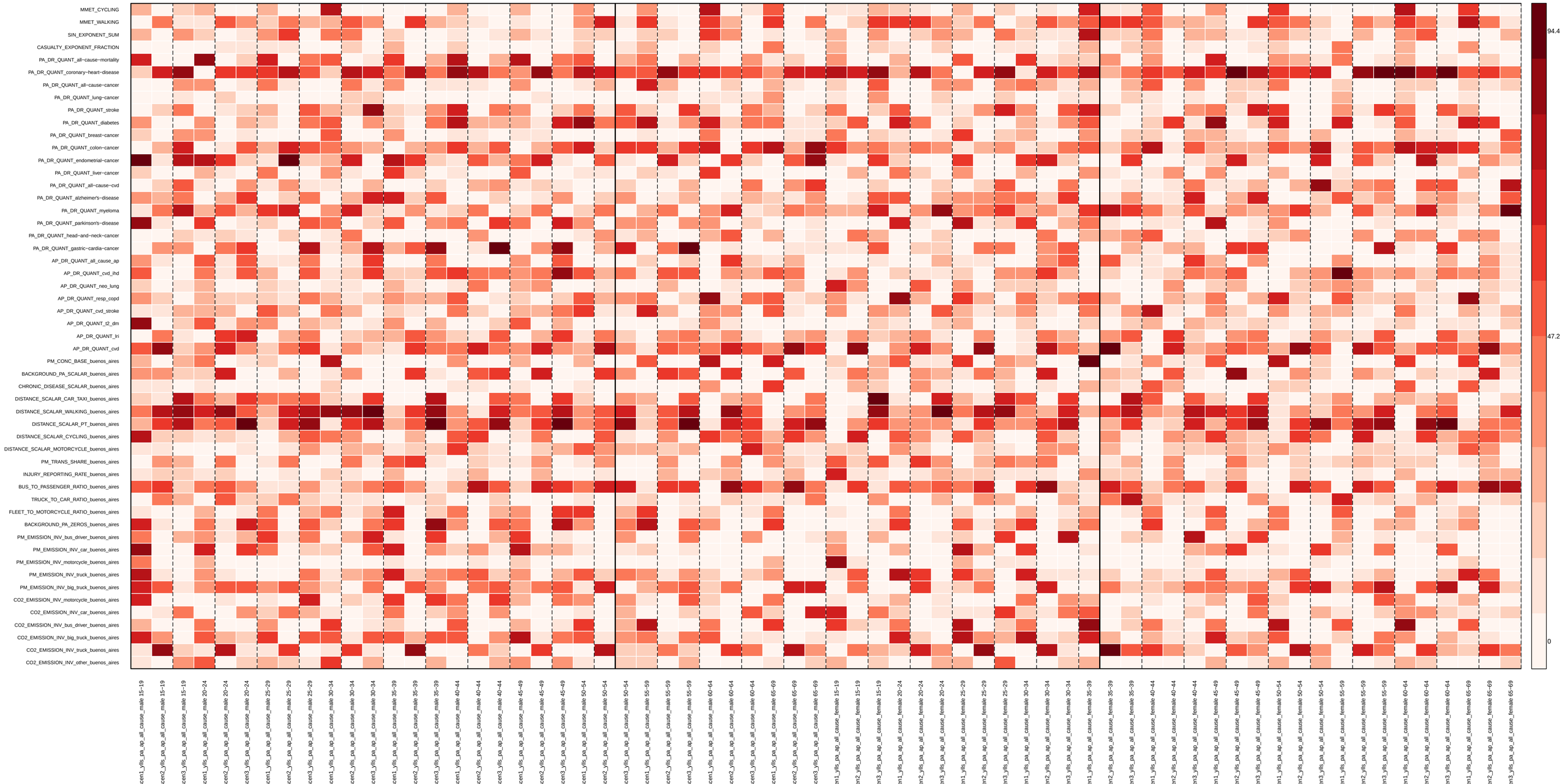
bogota − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



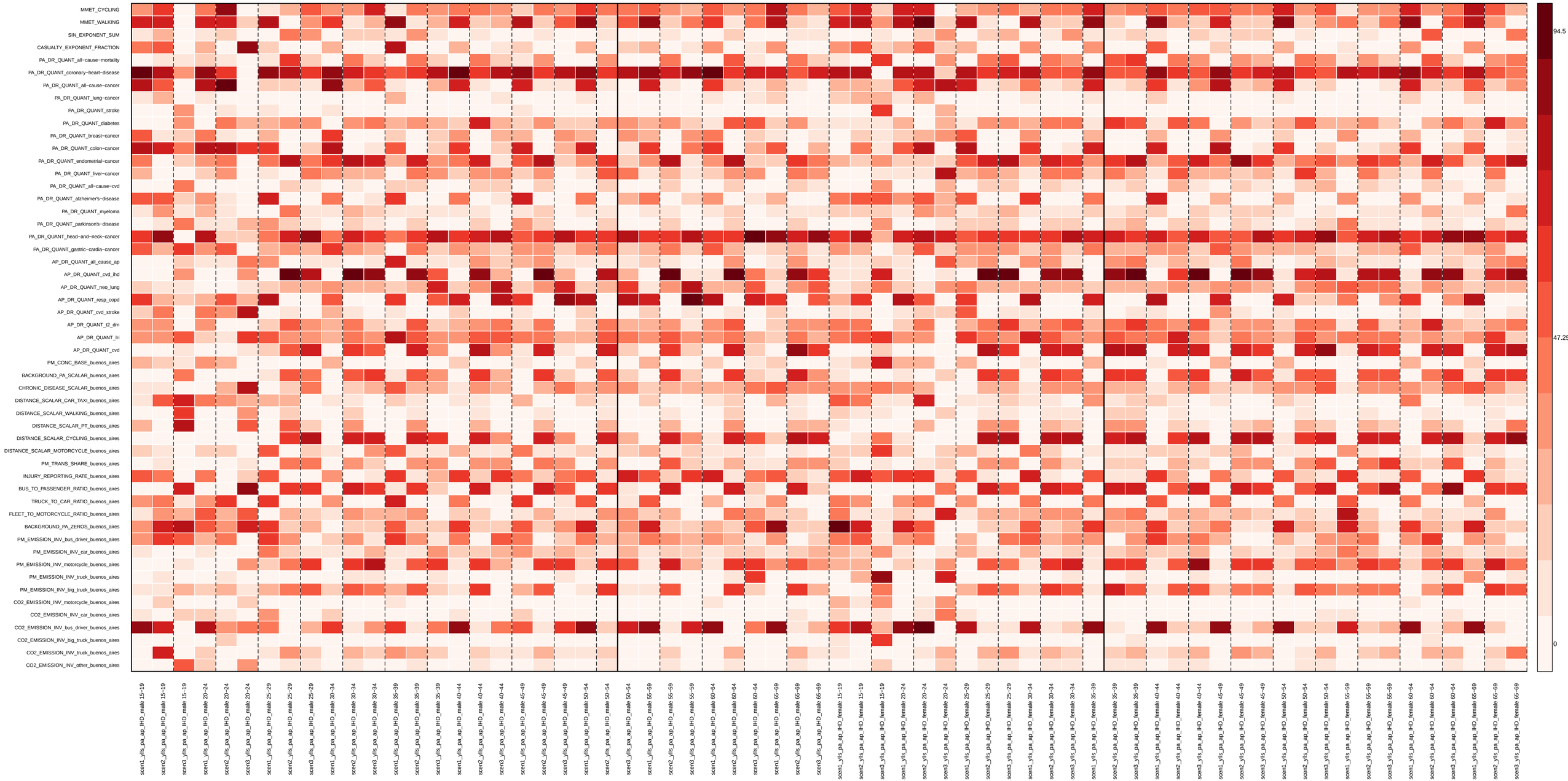
bogota − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



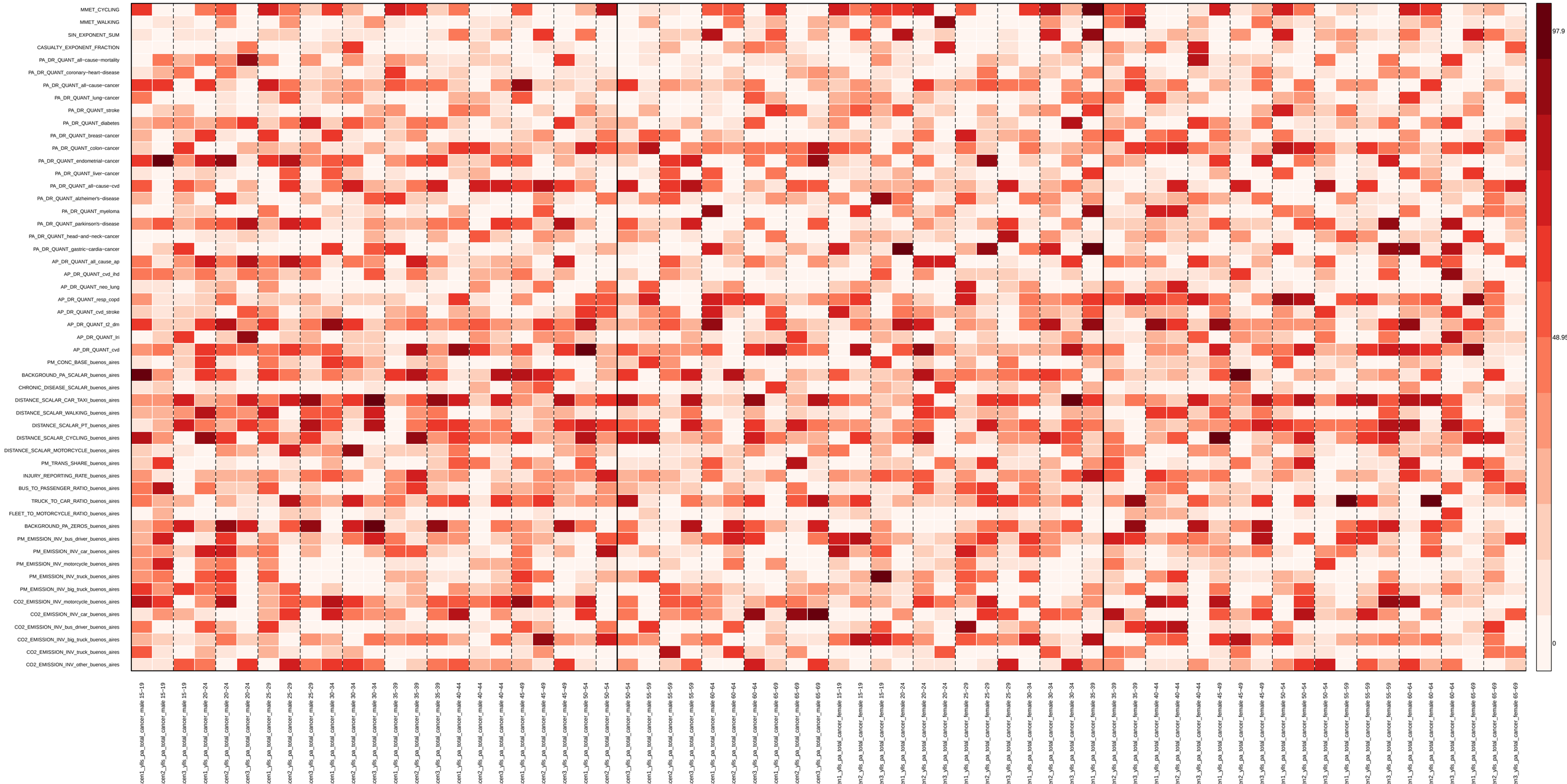
buenos_aires − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



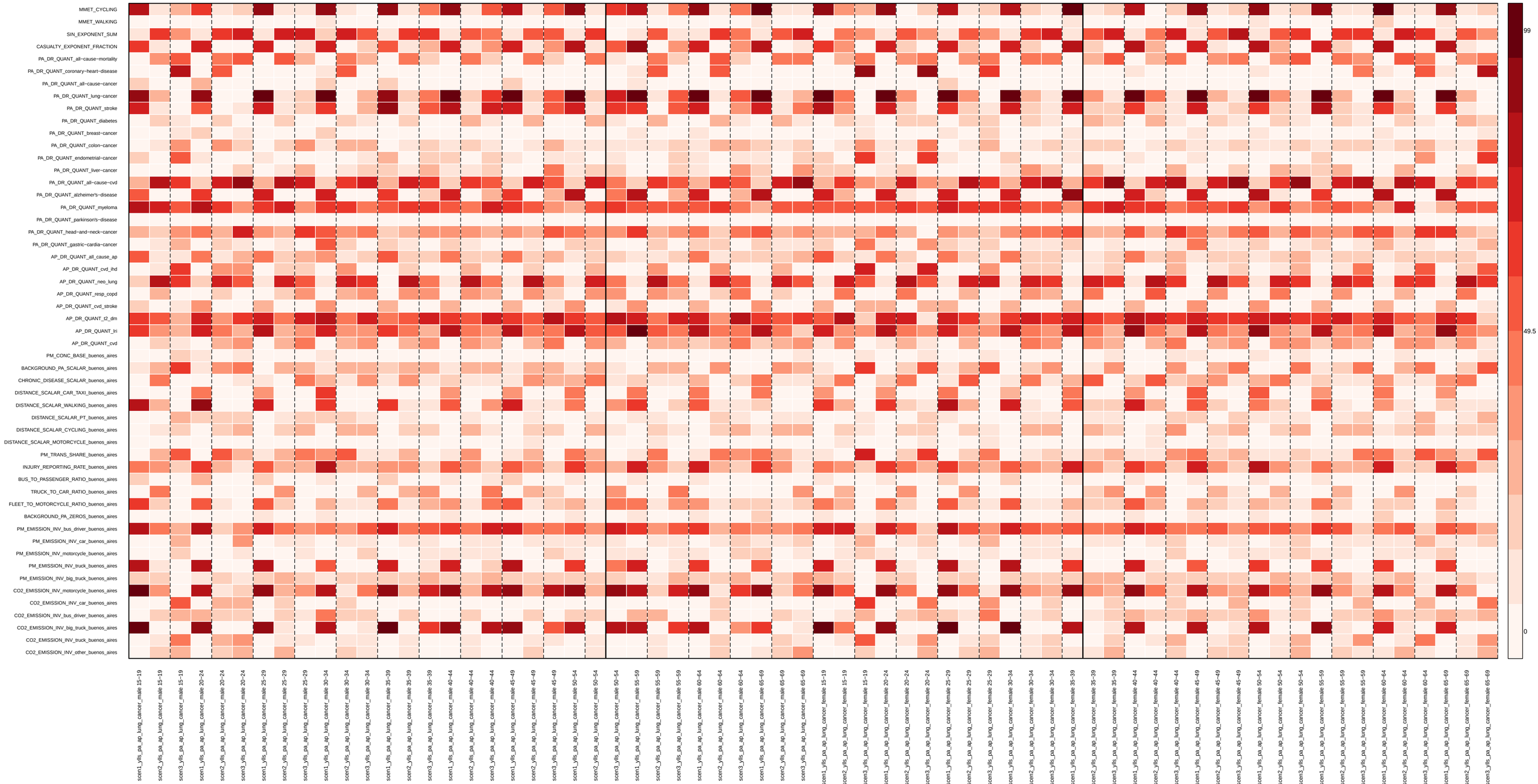
buenos_aires − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



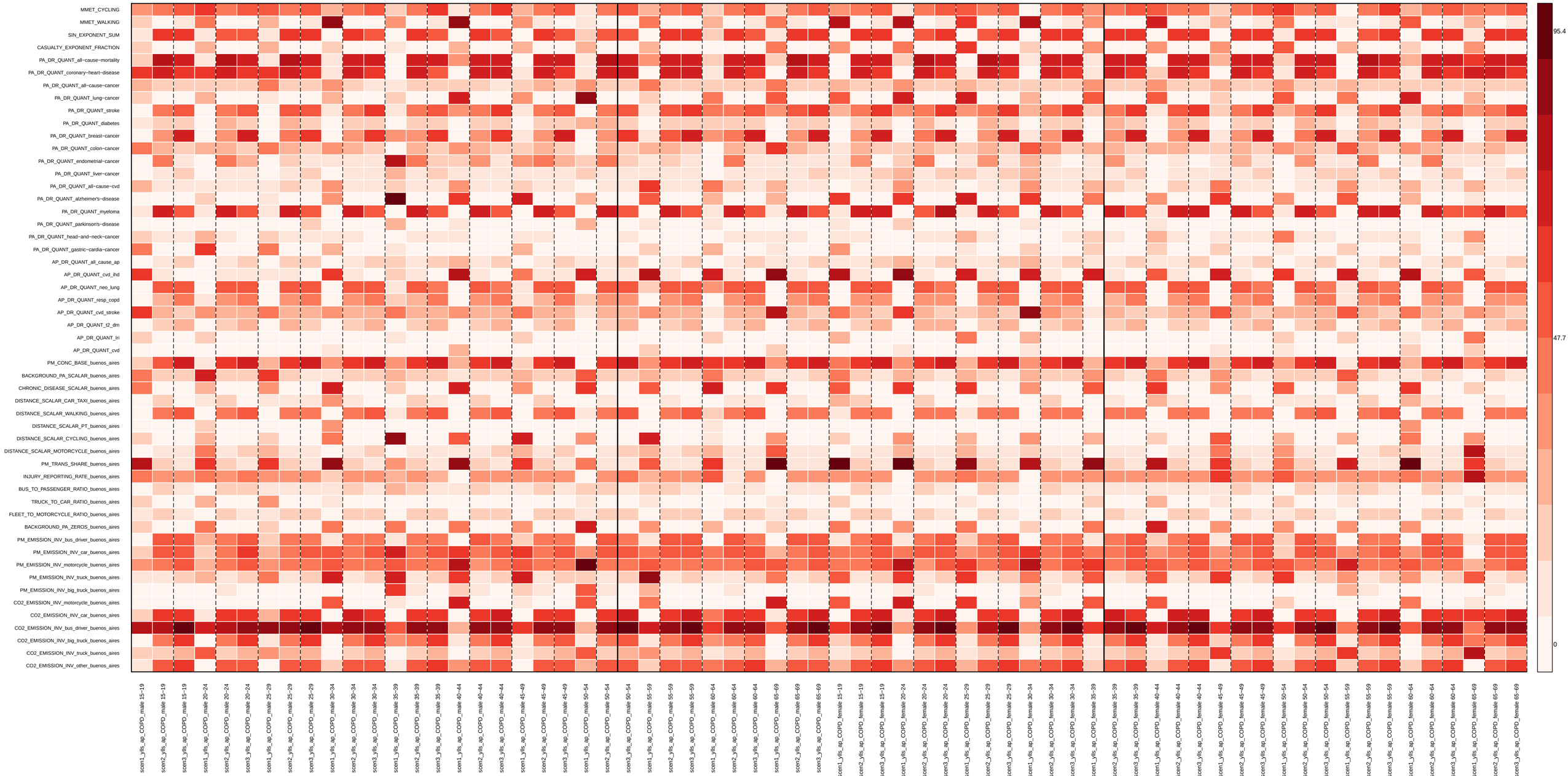
buenos_aires − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



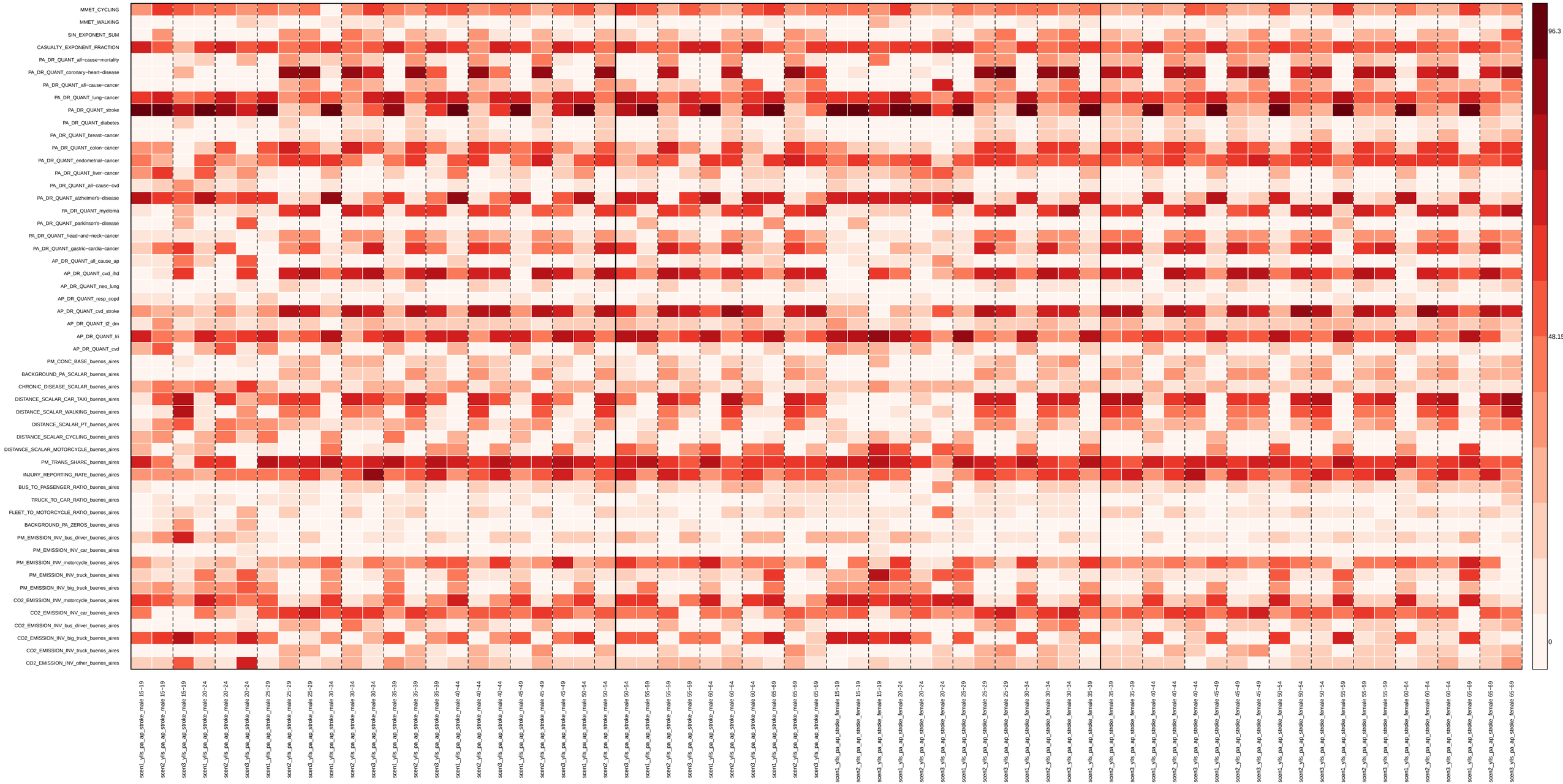
buenos_aires − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



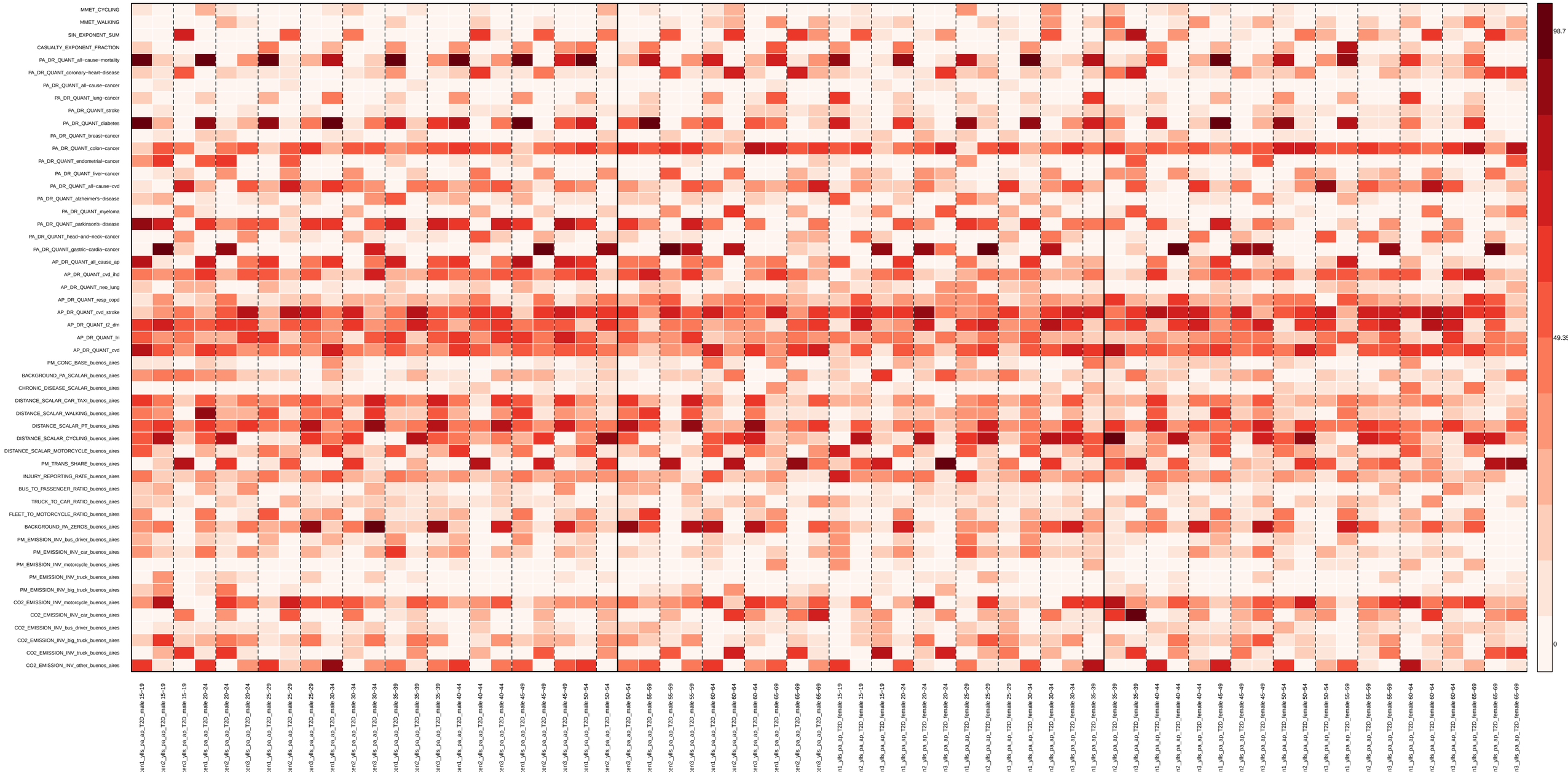
buenos_aires − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



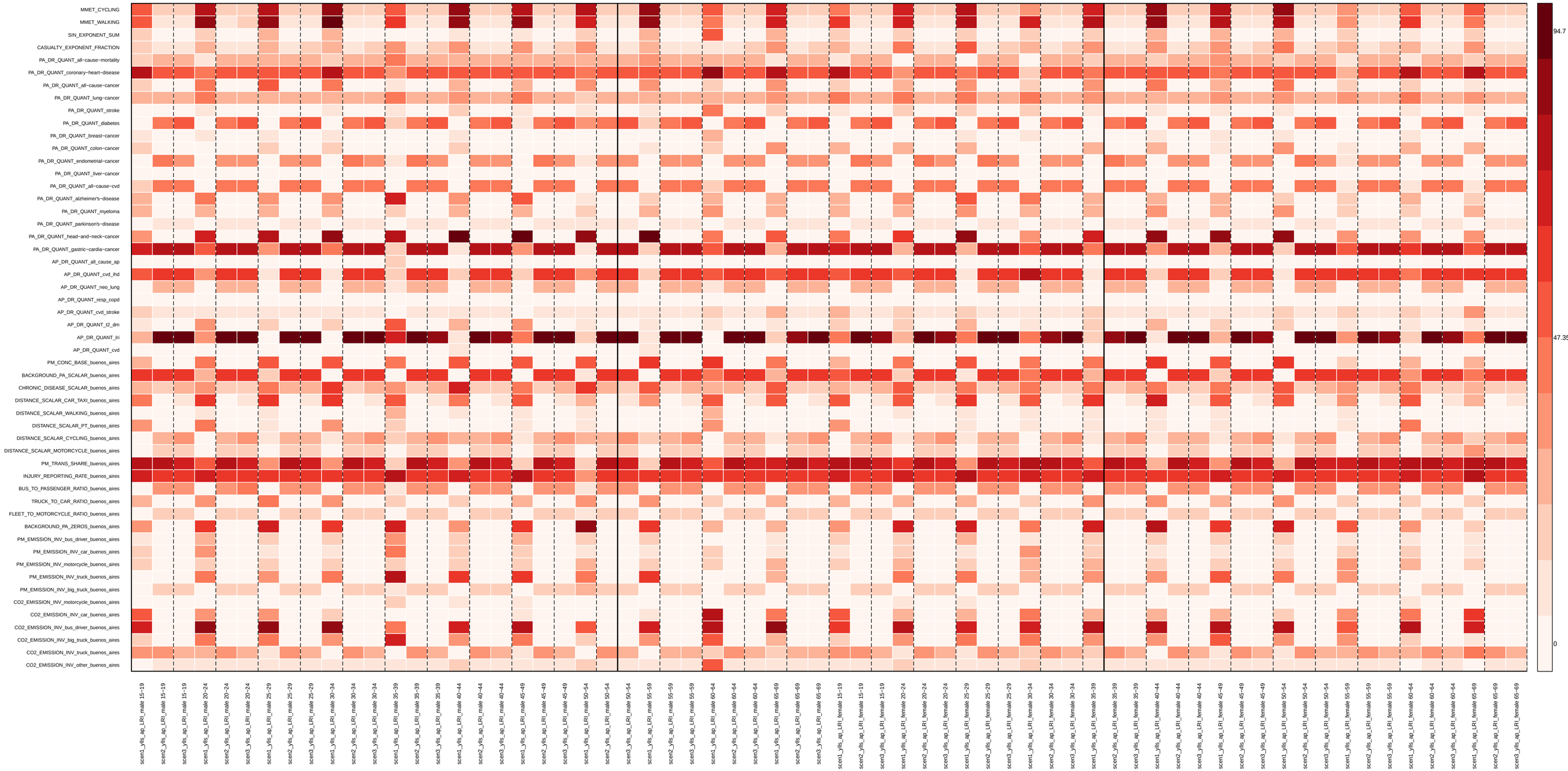
buenos_aires − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



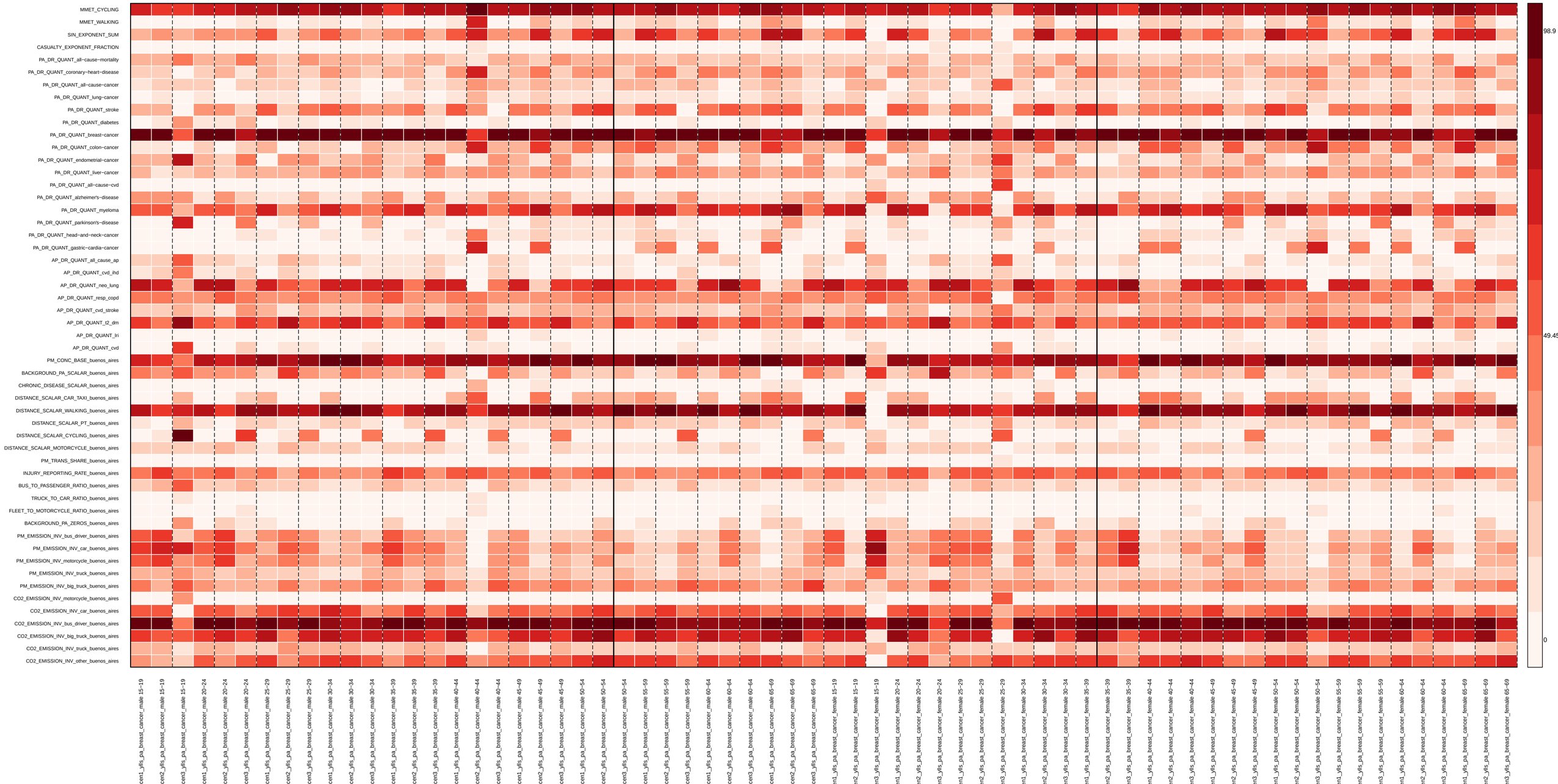
buenos_aires − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



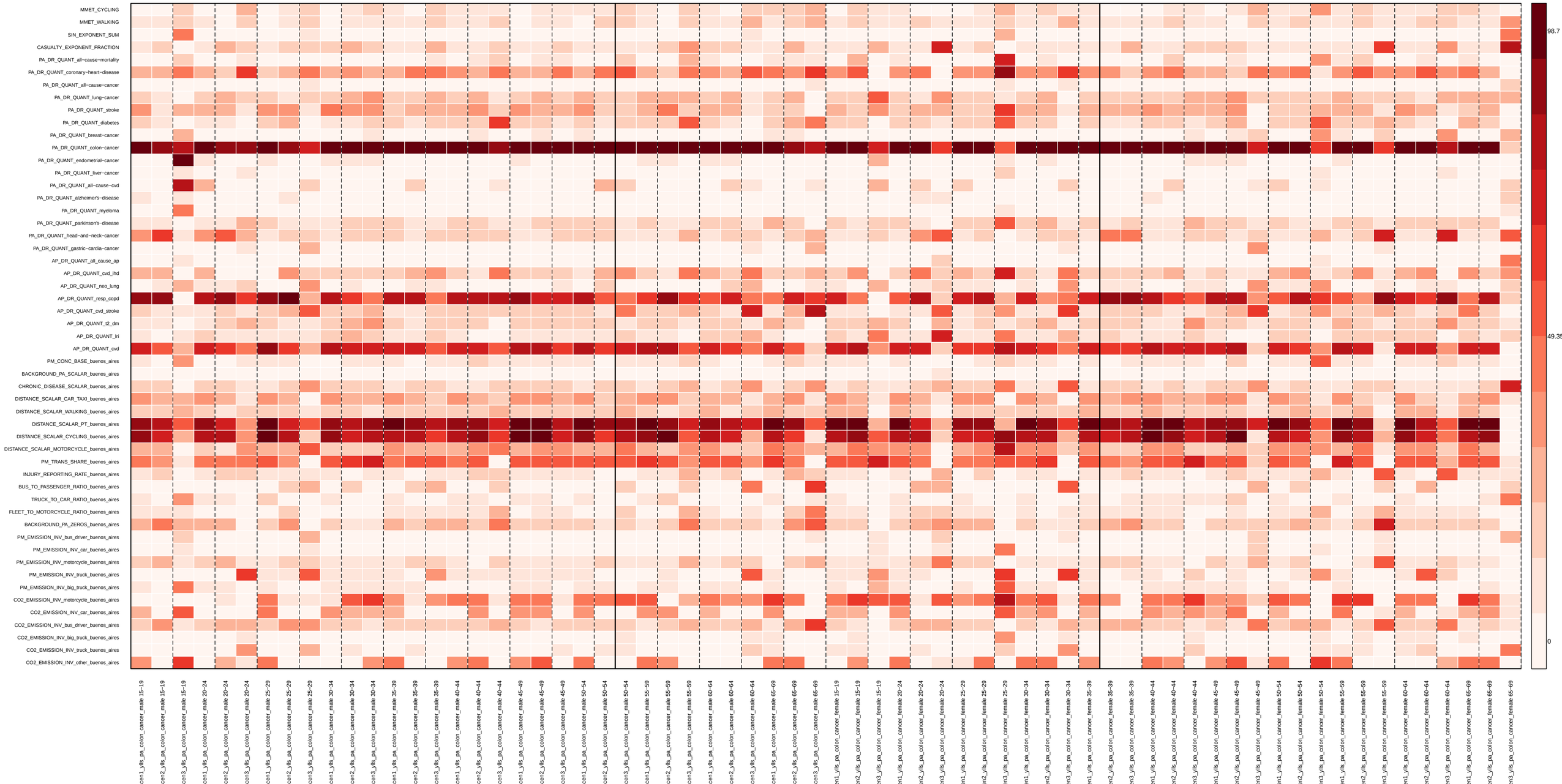
buenos_aires − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



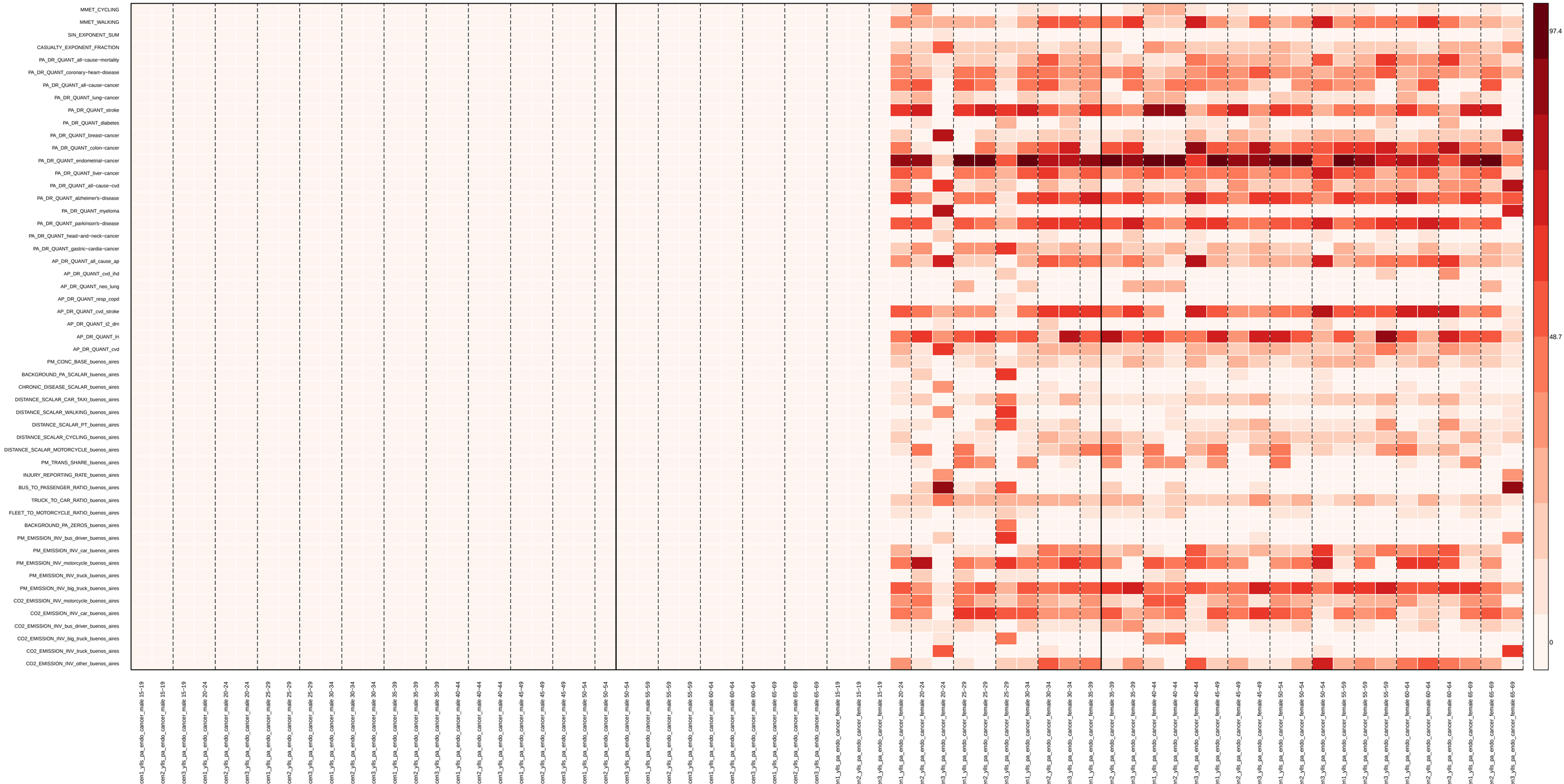
buenos_aires − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



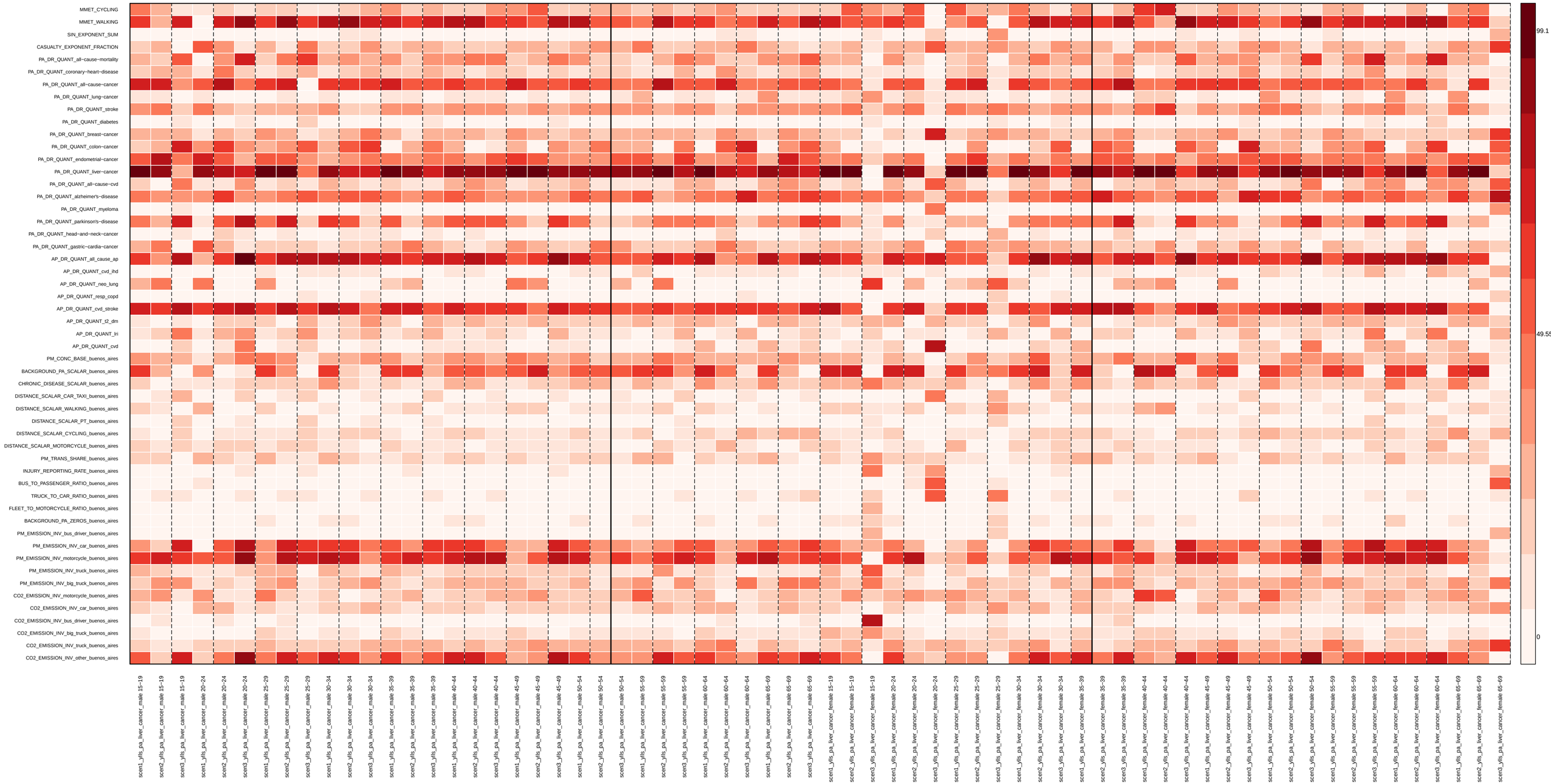
buenos_aires − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



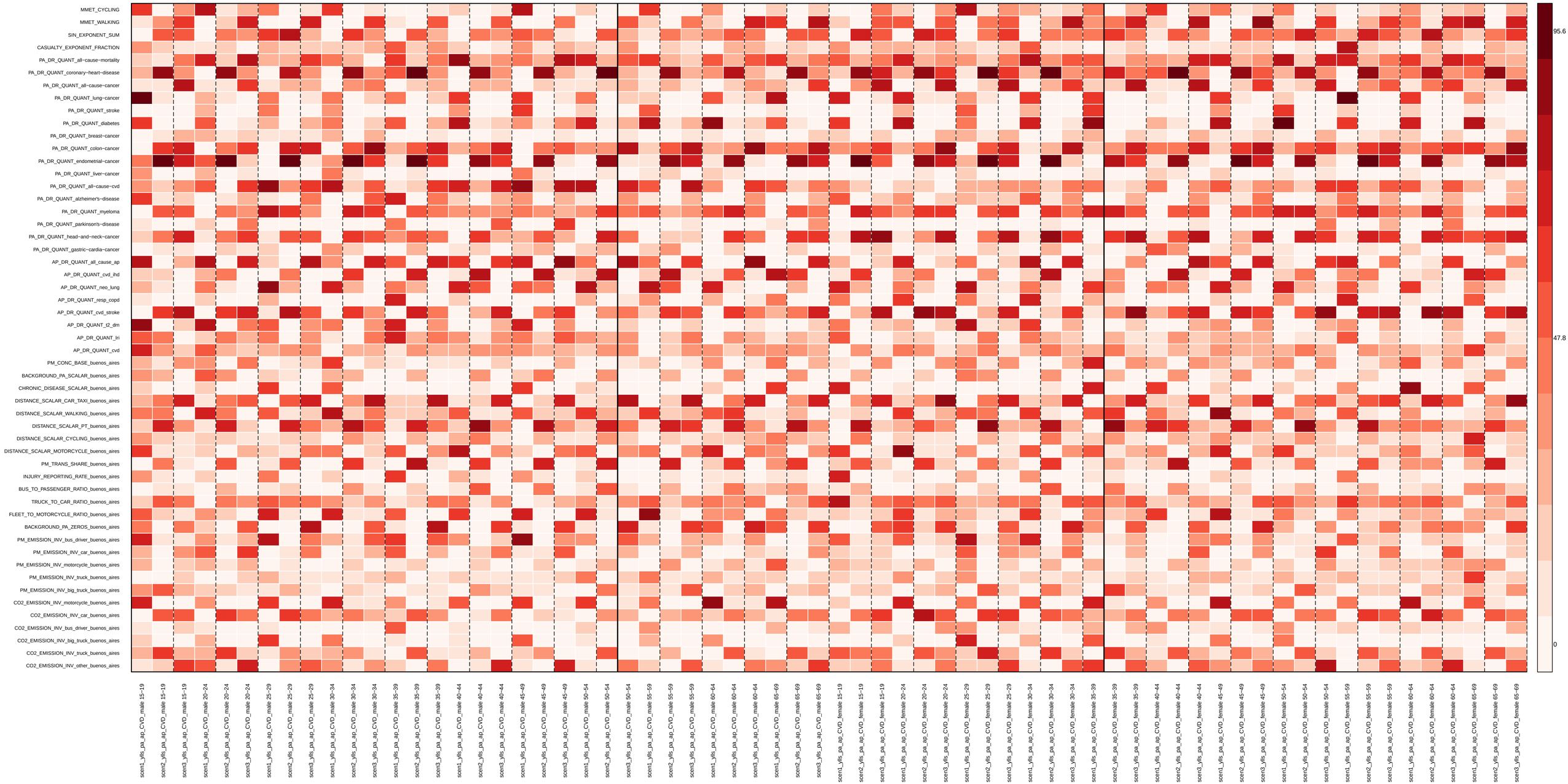
buenos_aires − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



buenos_aires − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

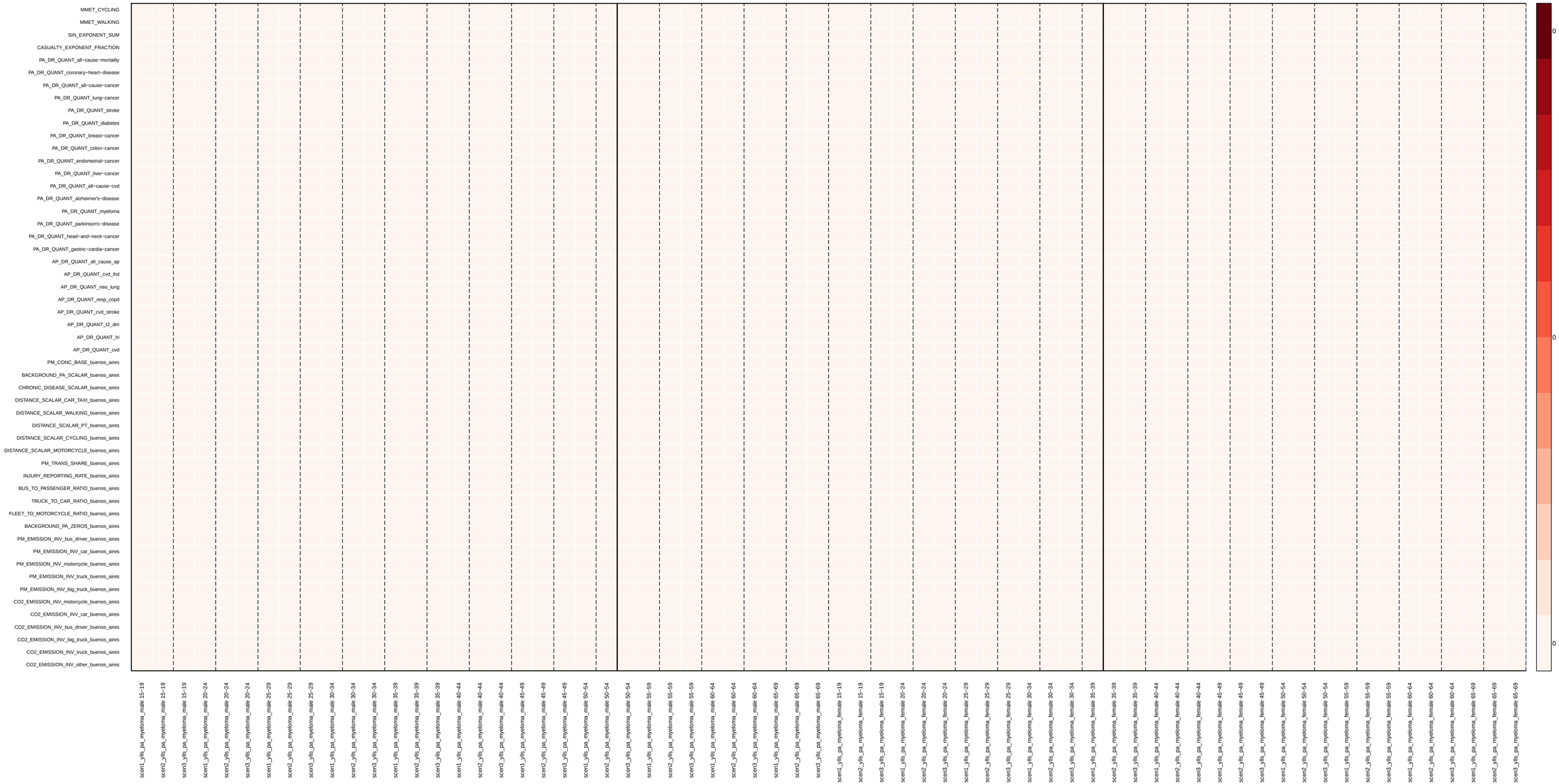


buenos_aires − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

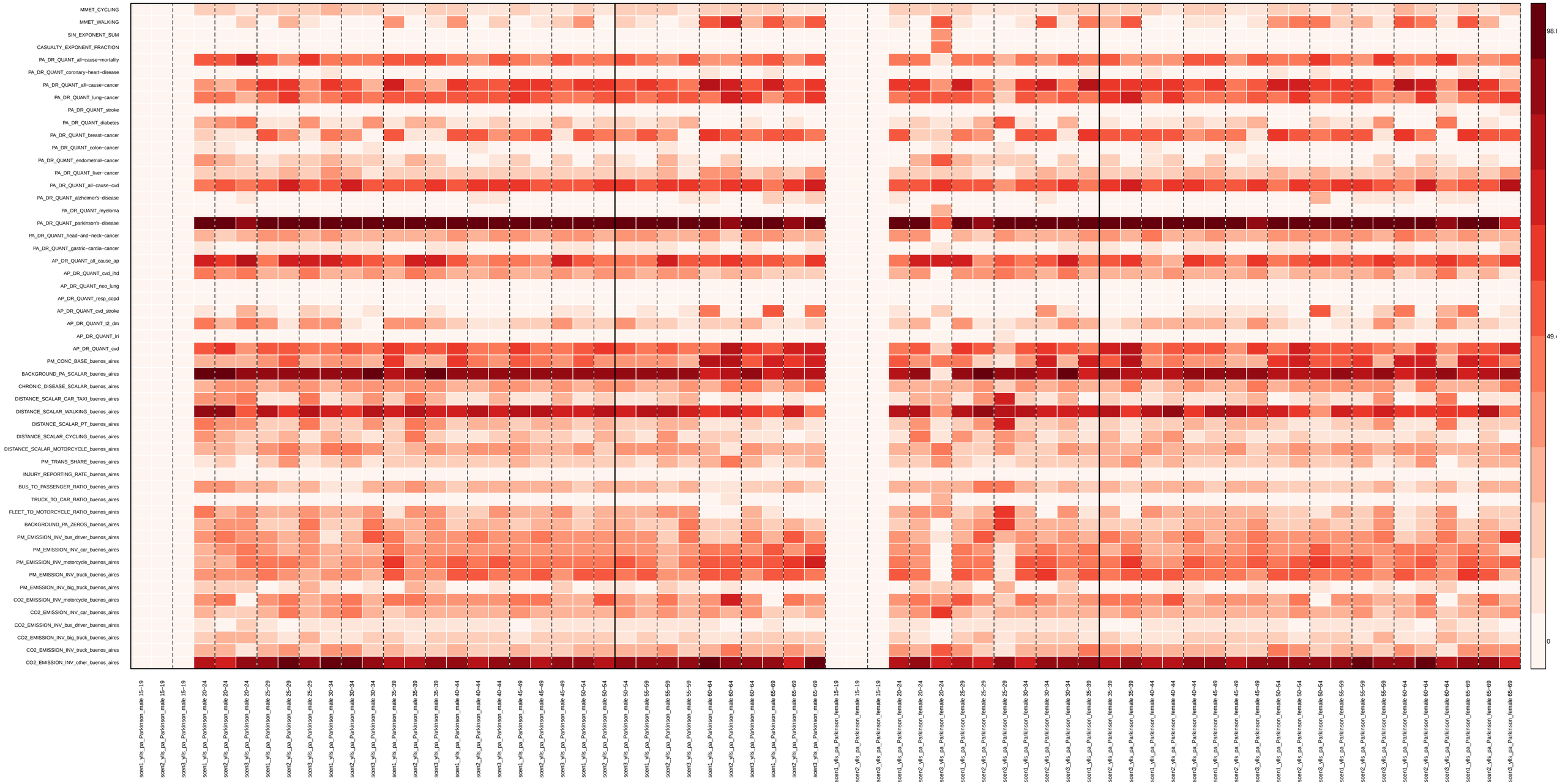


buenos_aires − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

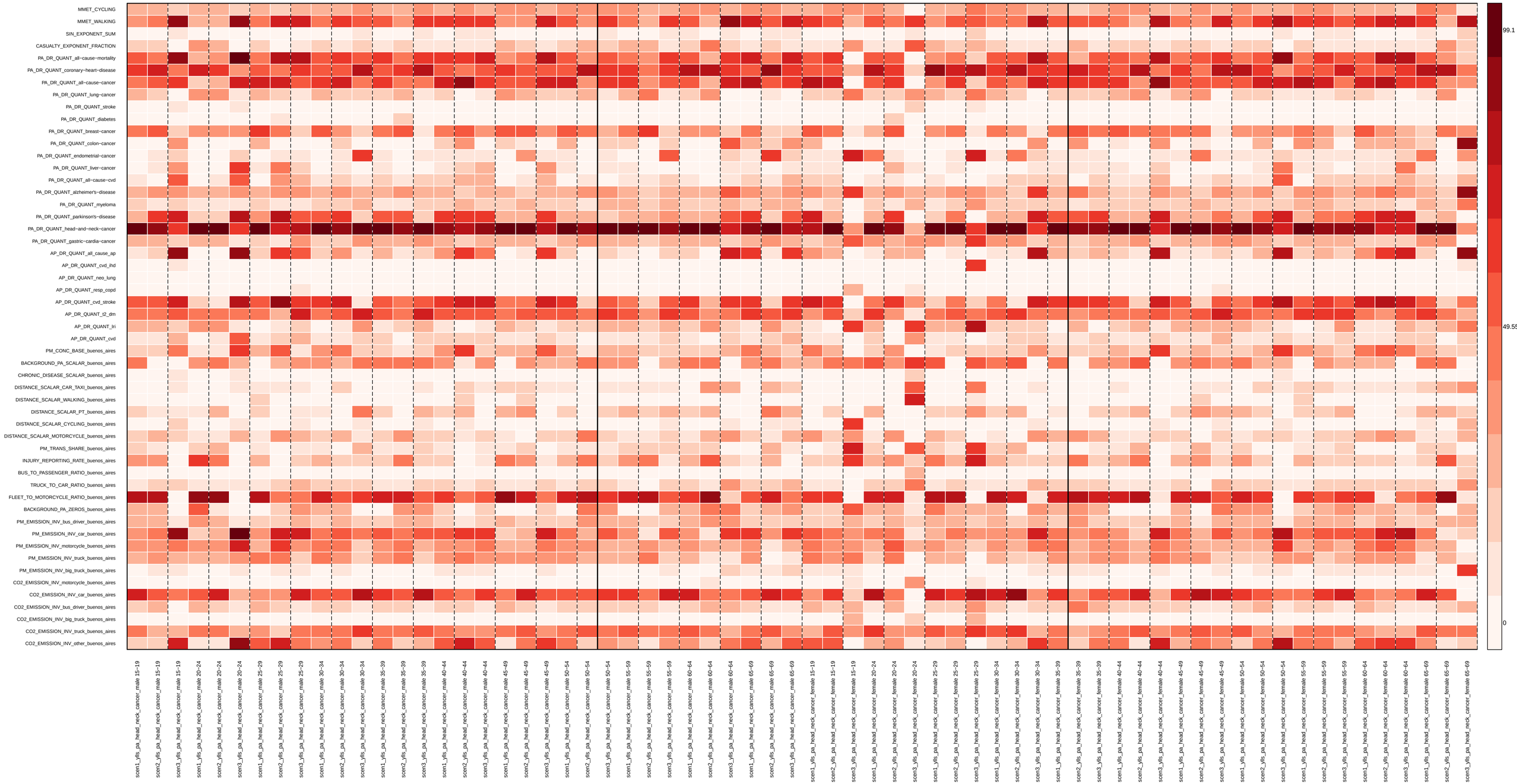
buenos_aires − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



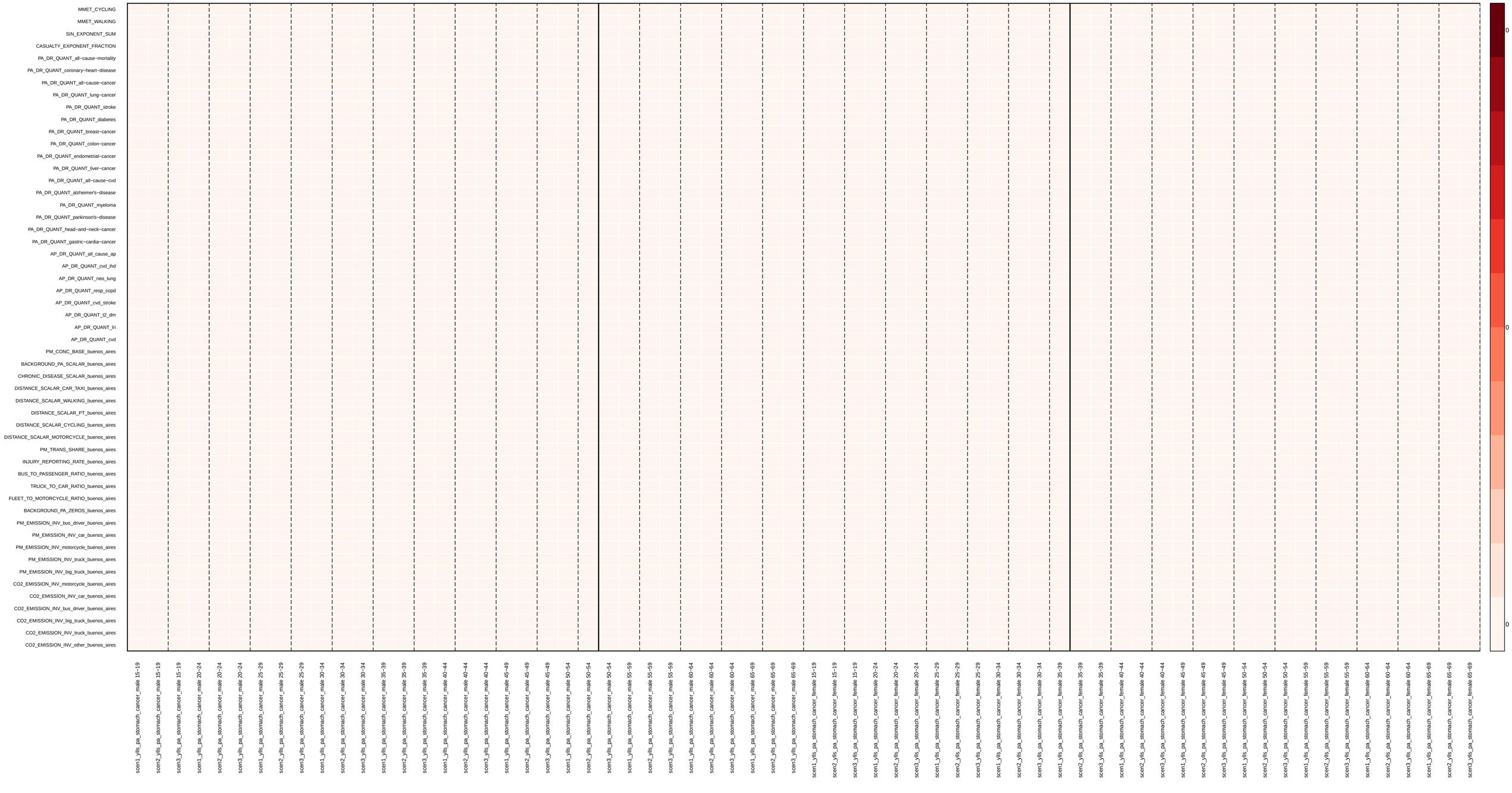
buenos_aires − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



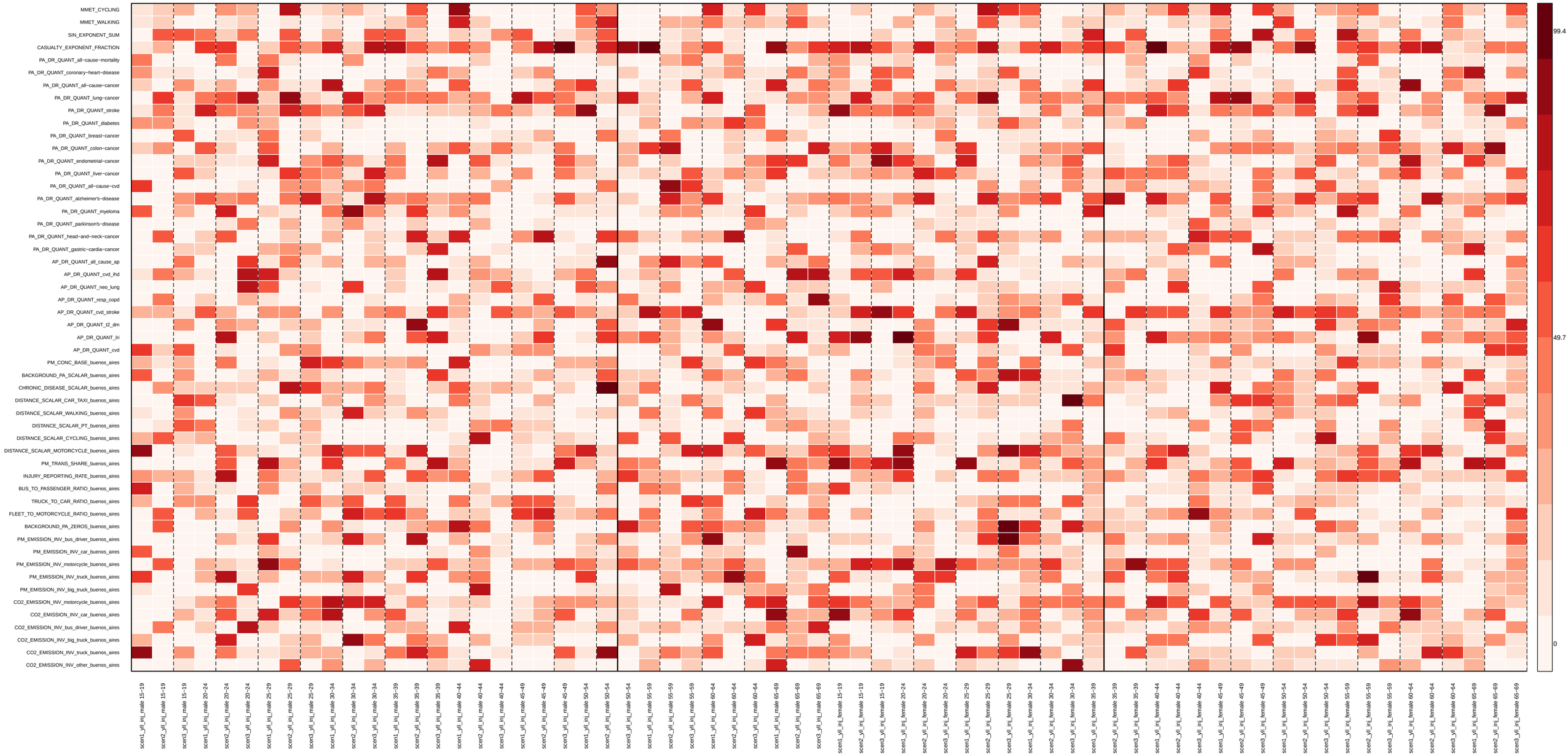
buenos_aires − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



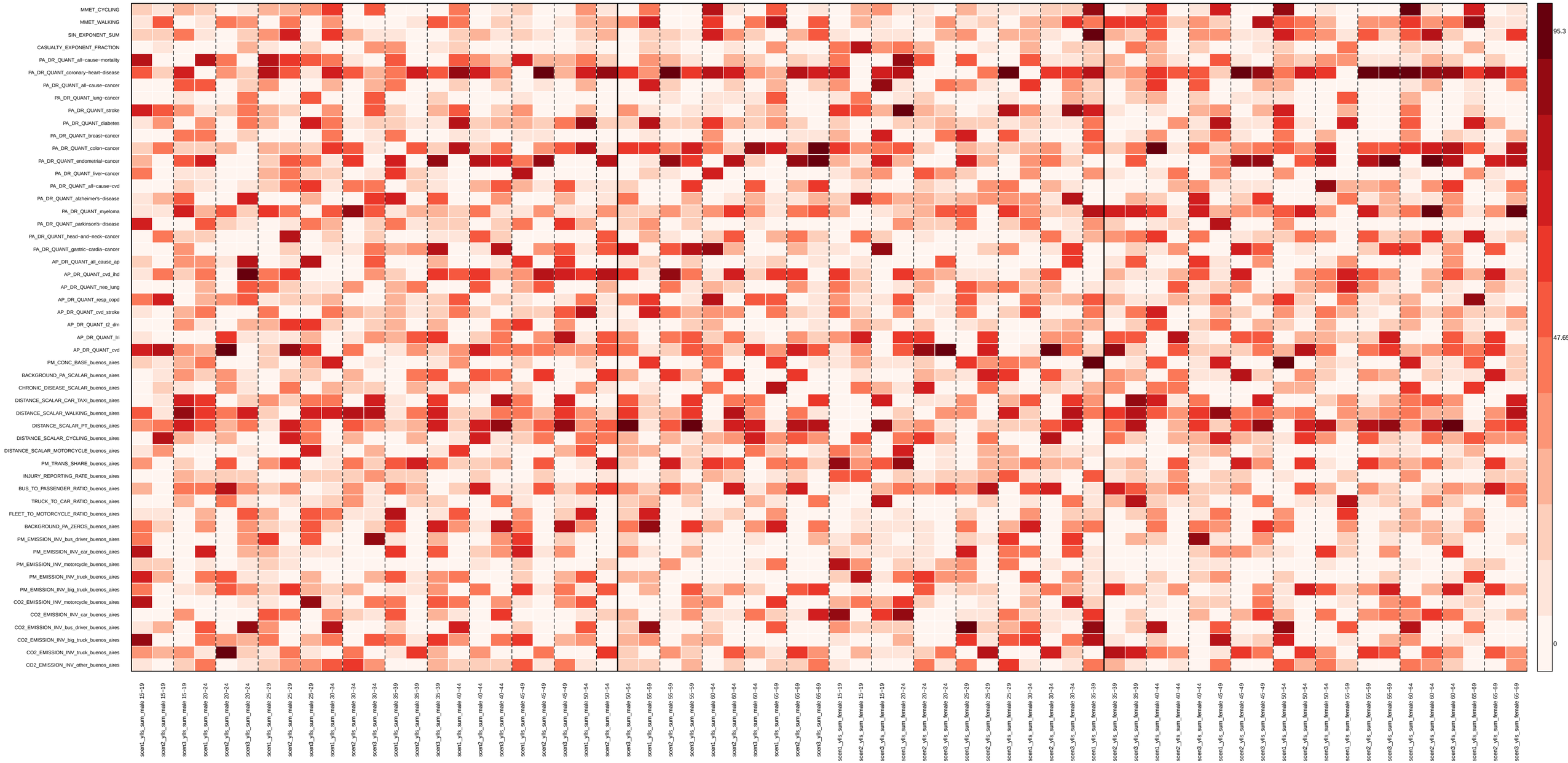
buenos_aires − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



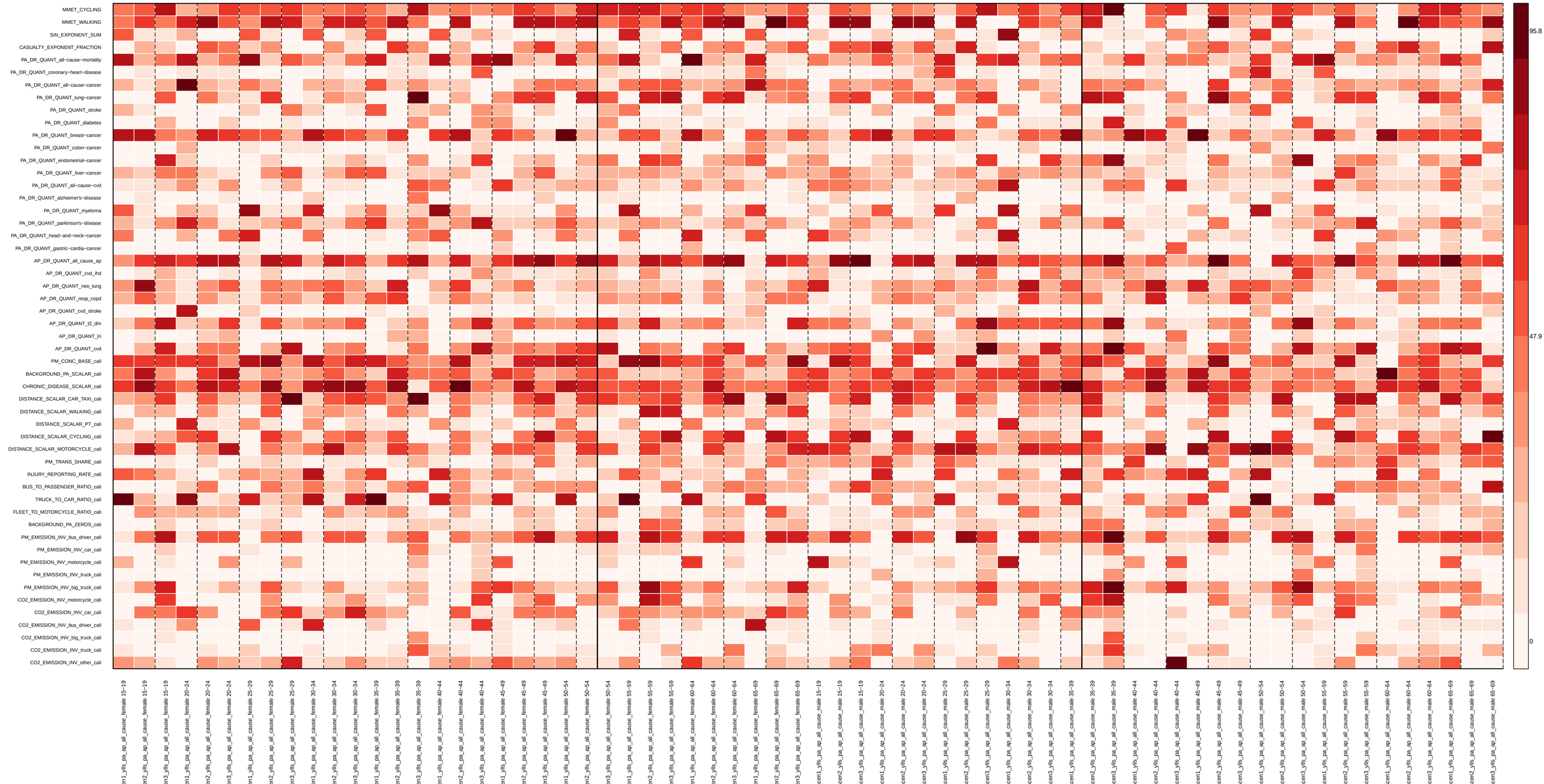
buenos_aires − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



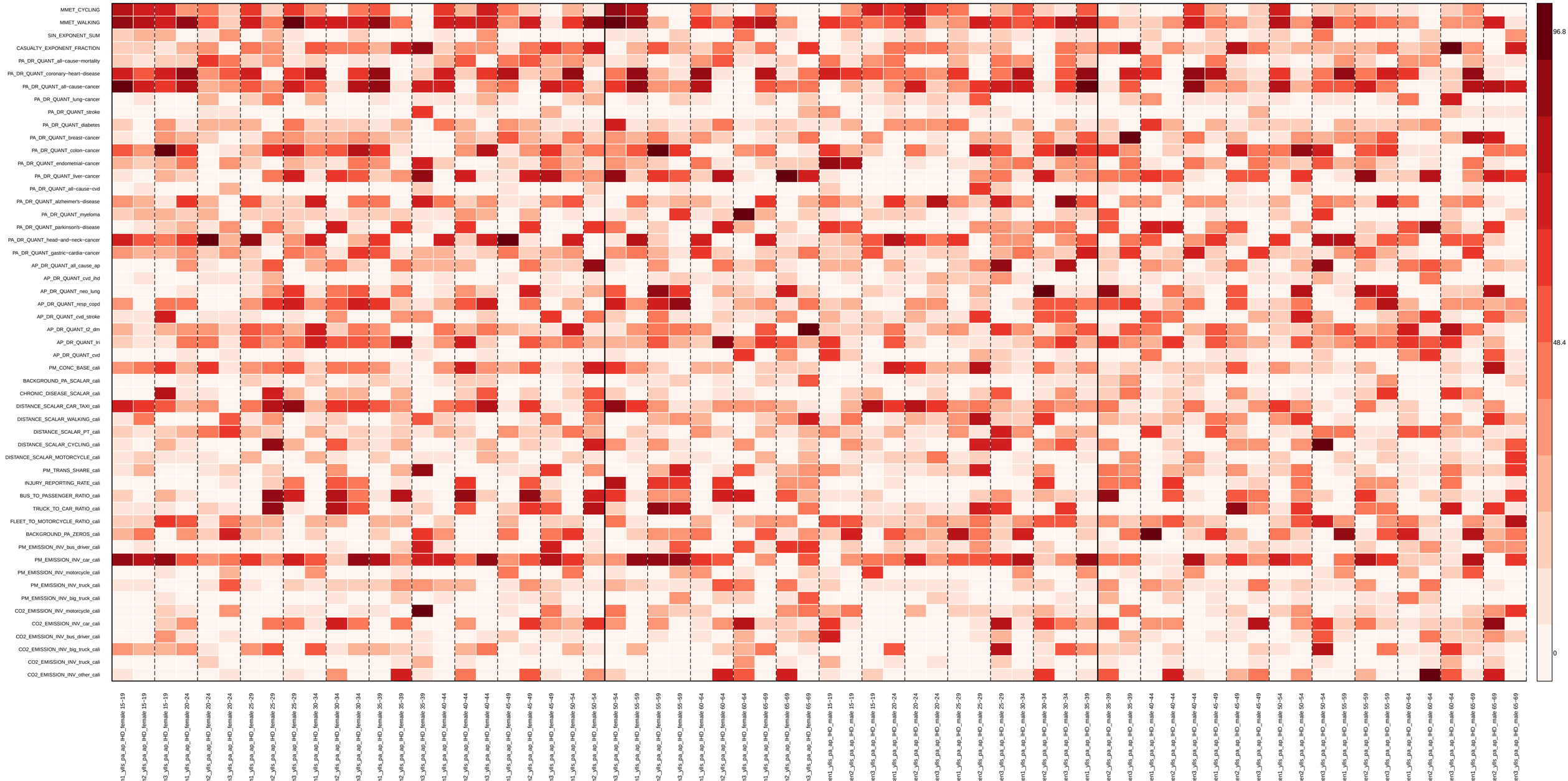
buenos_aires − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



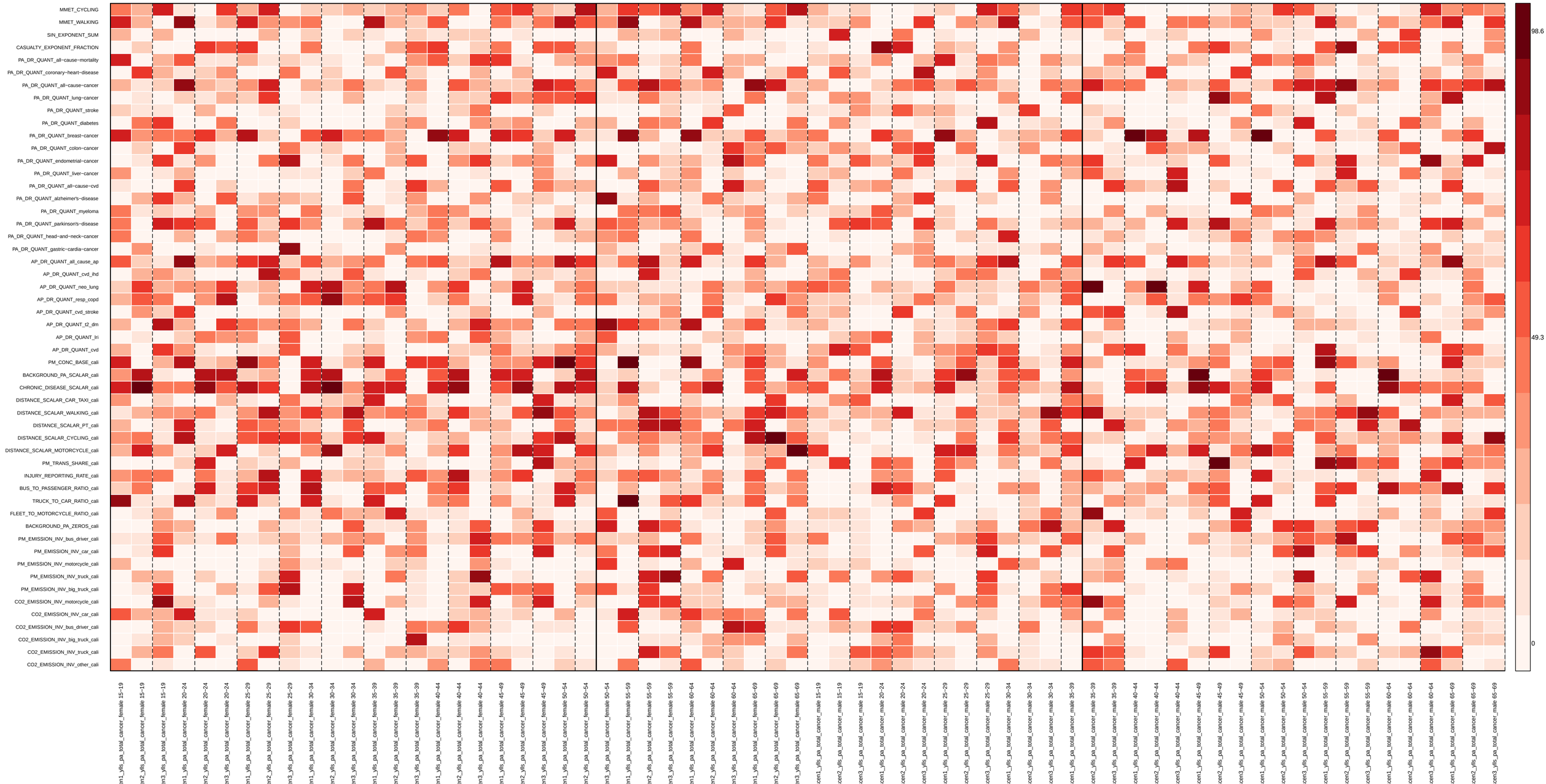
cali − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



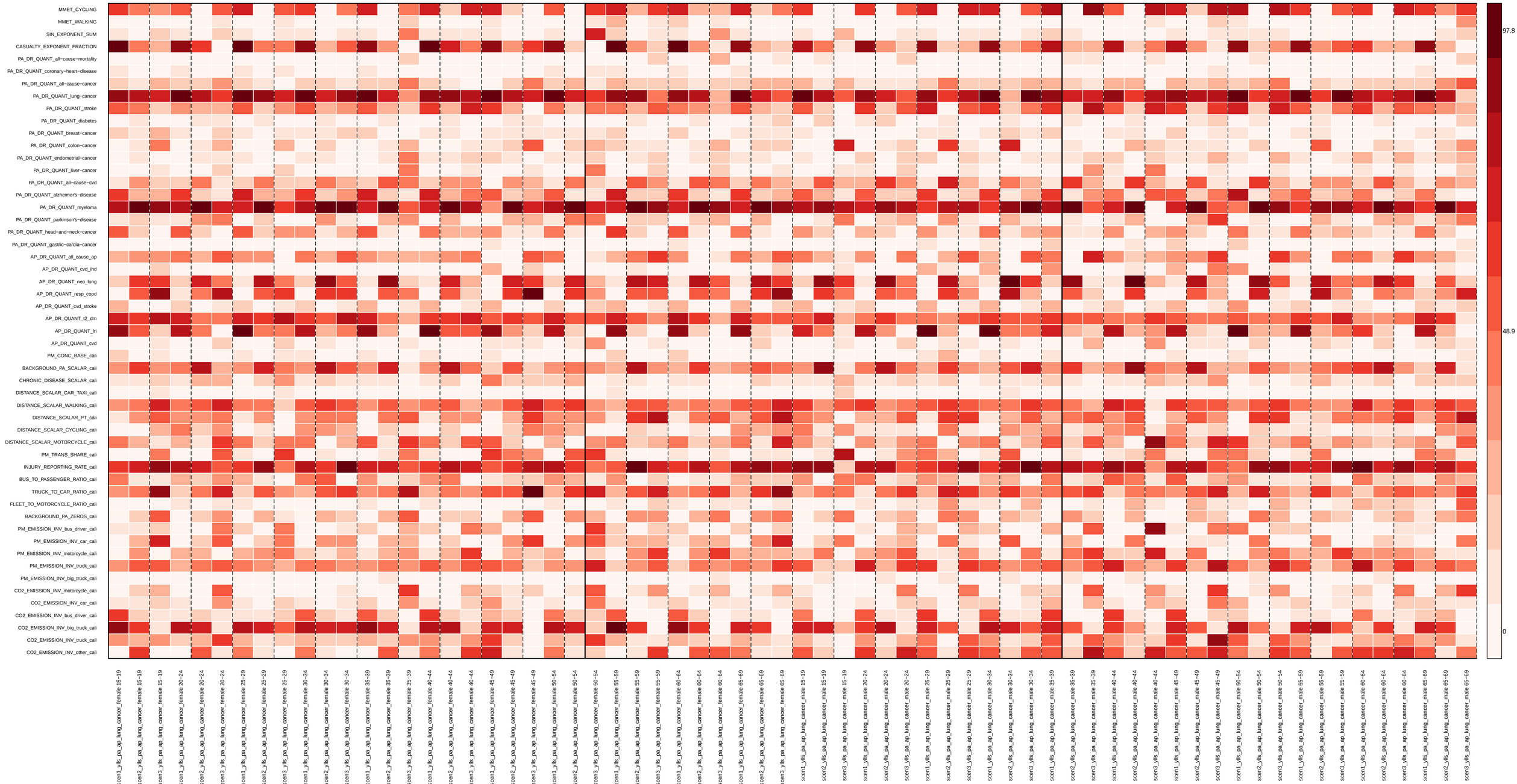
cali − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



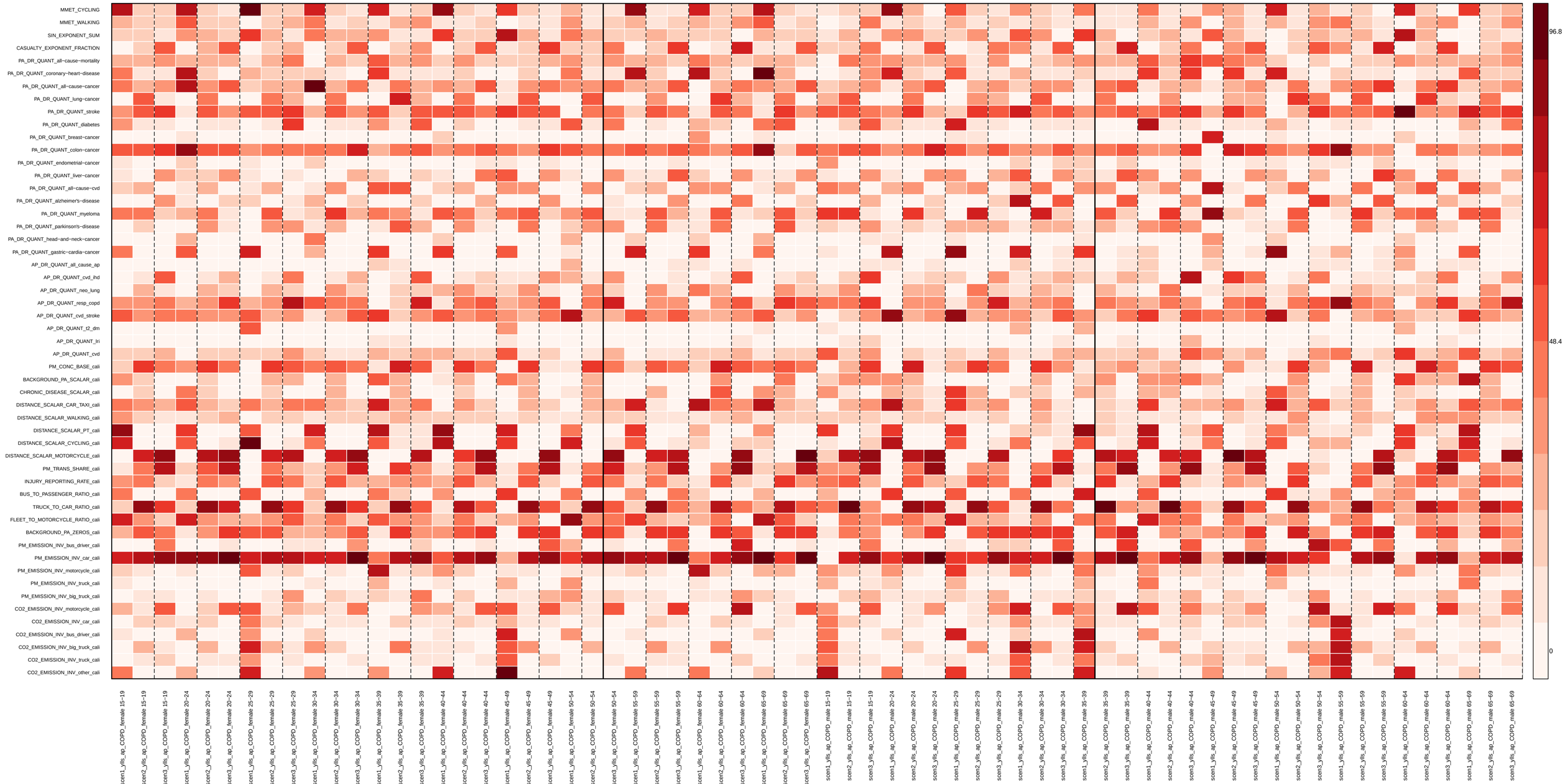
cali − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



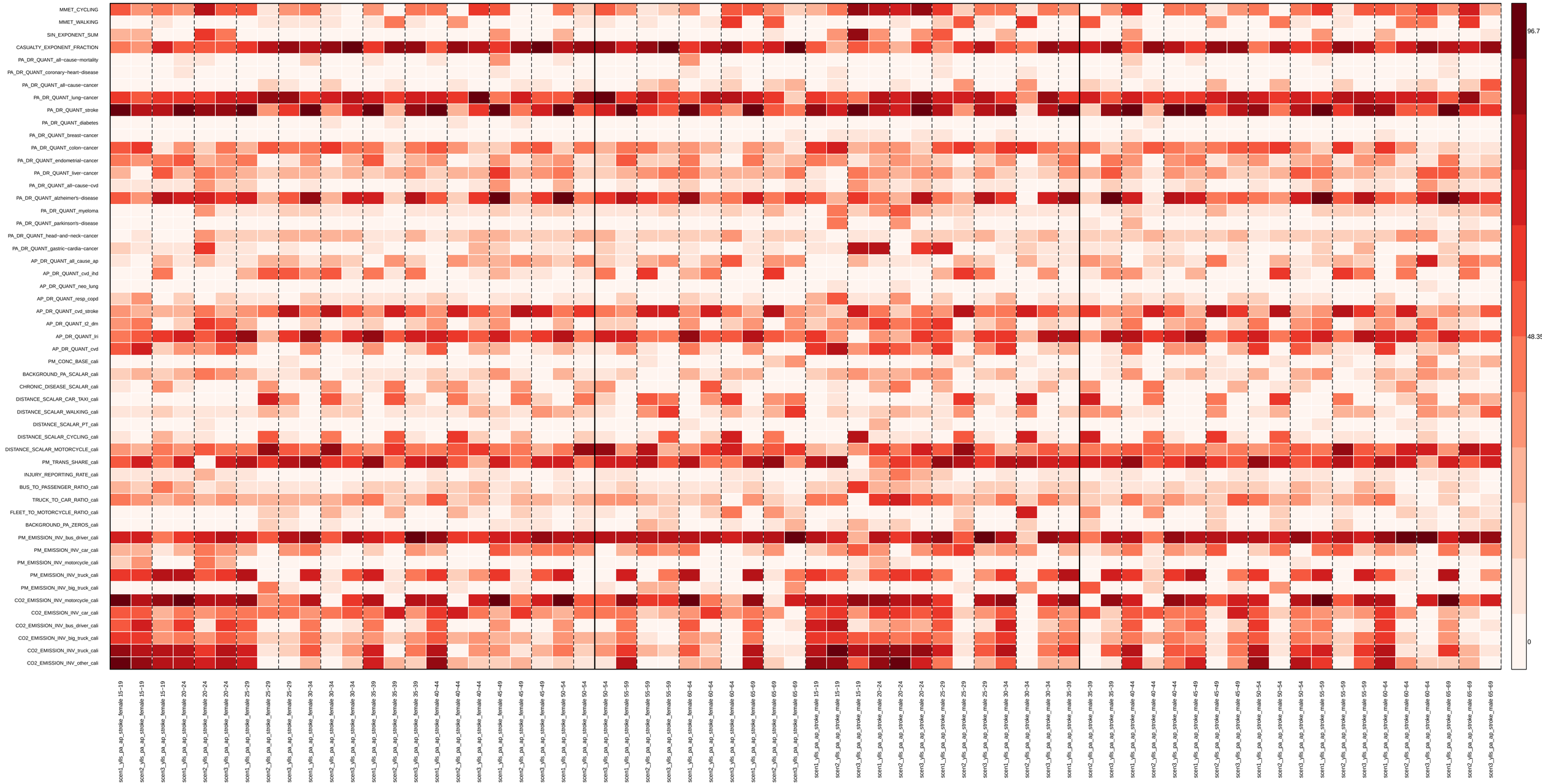
cali − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



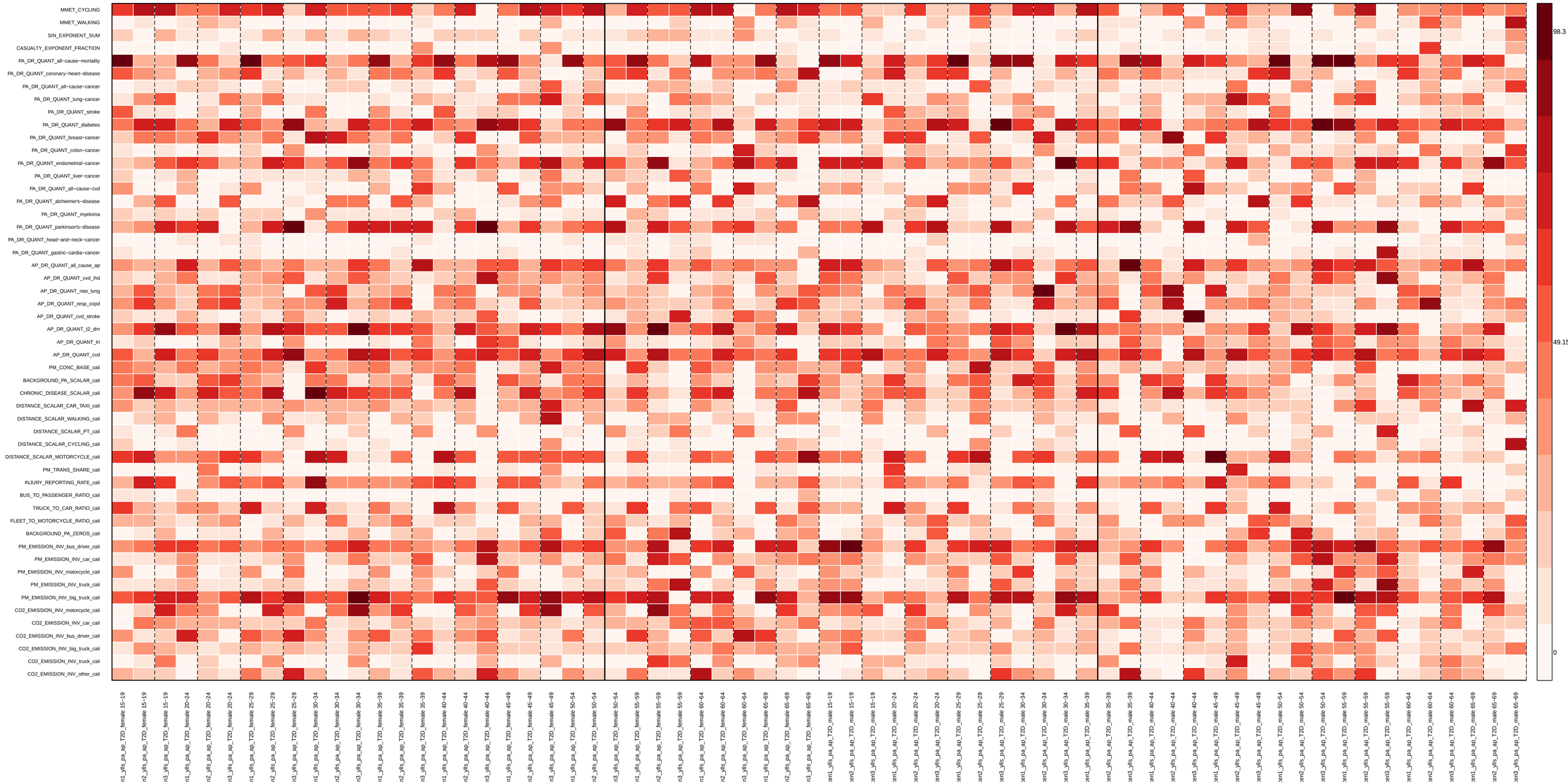
cali − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



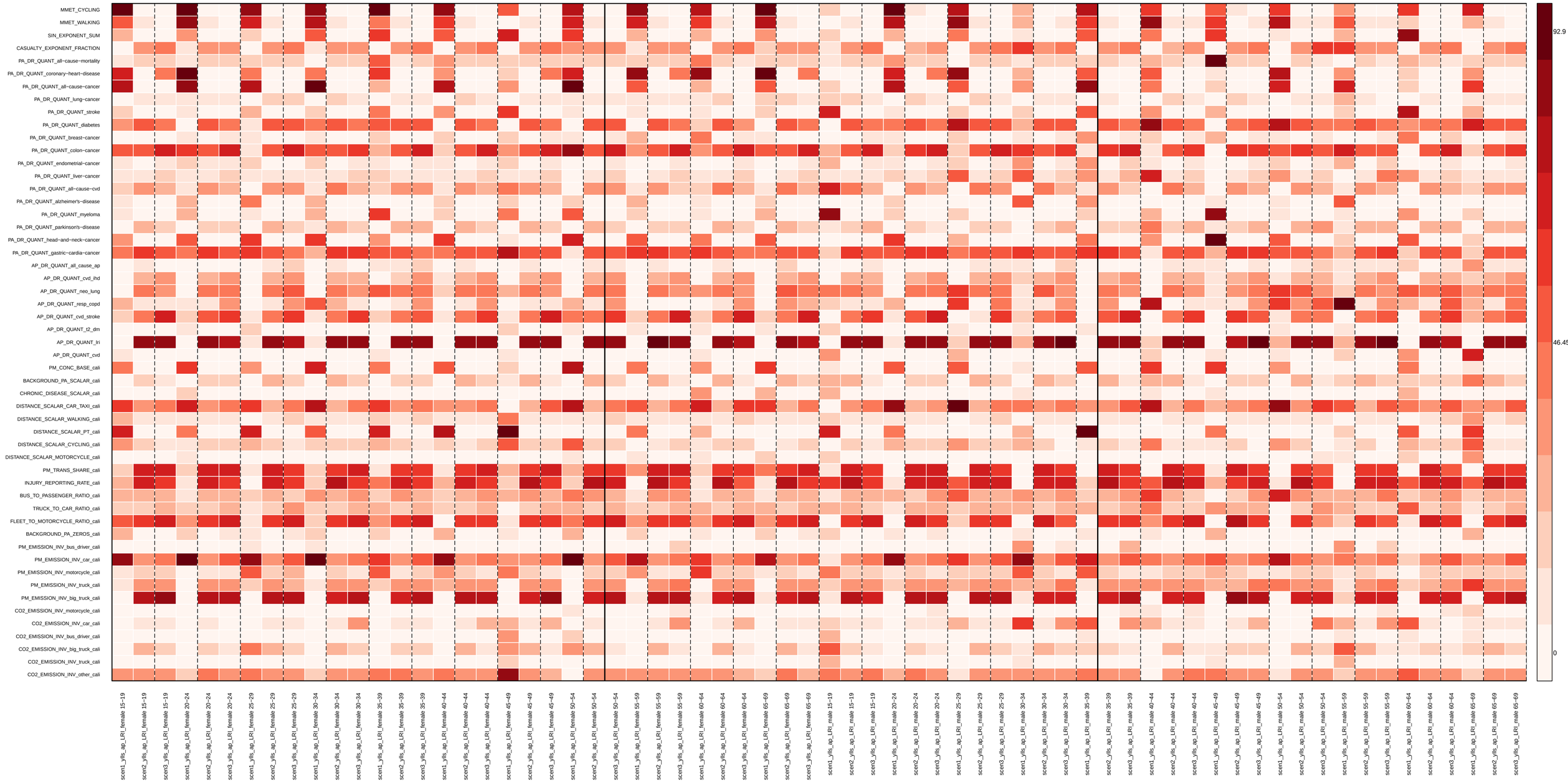
cali − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



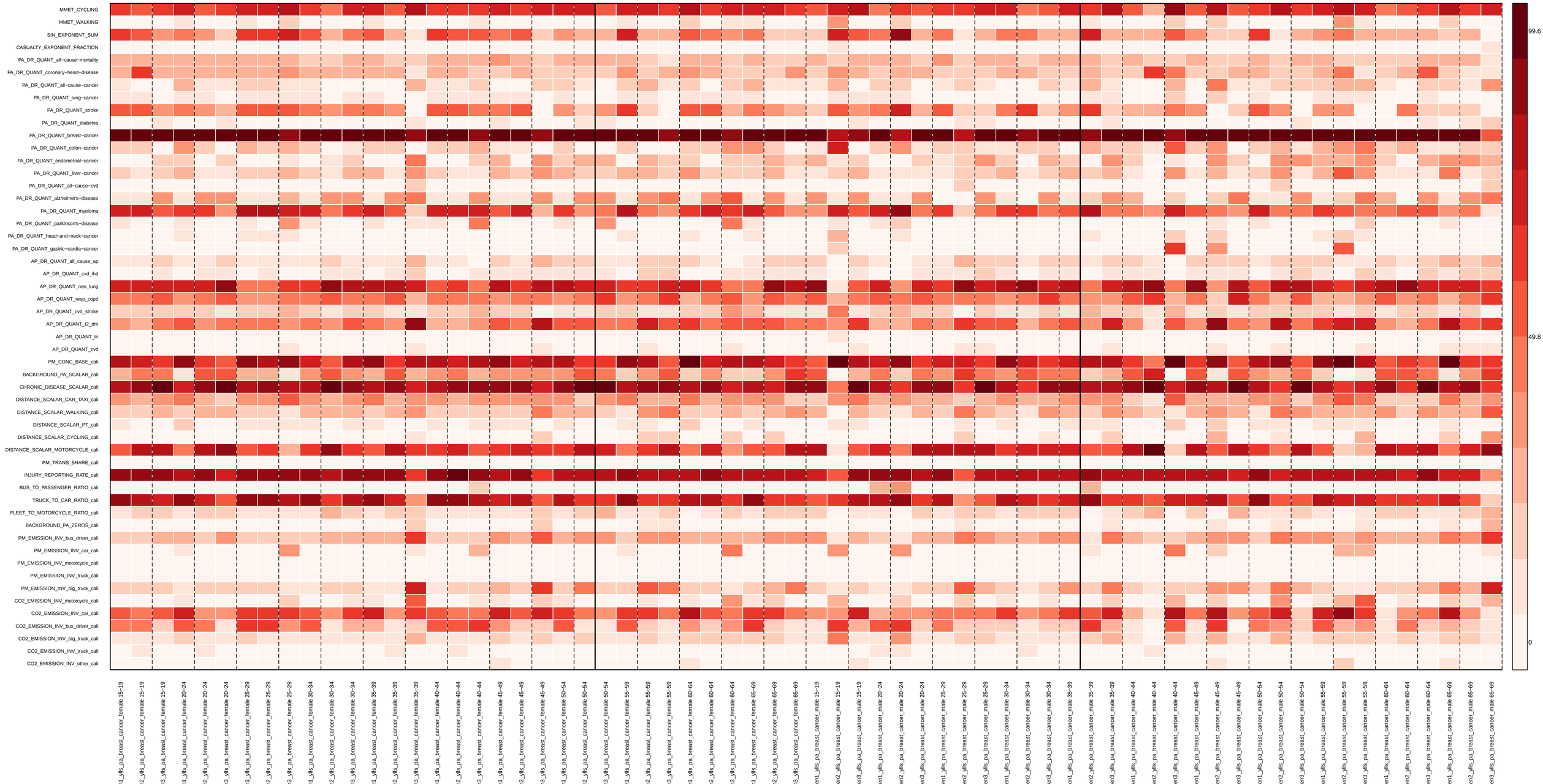
cali − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



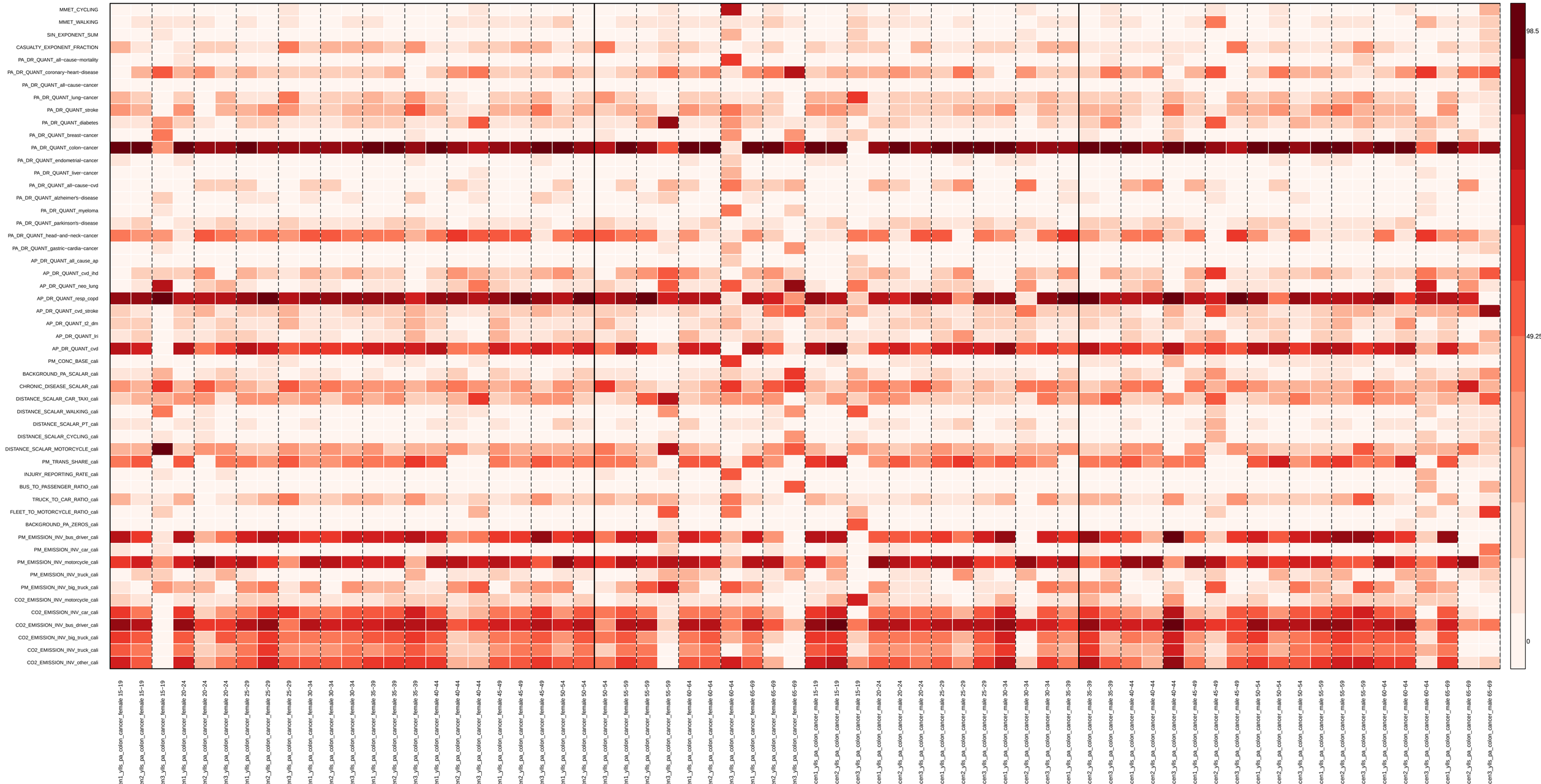
cali − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



cali − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

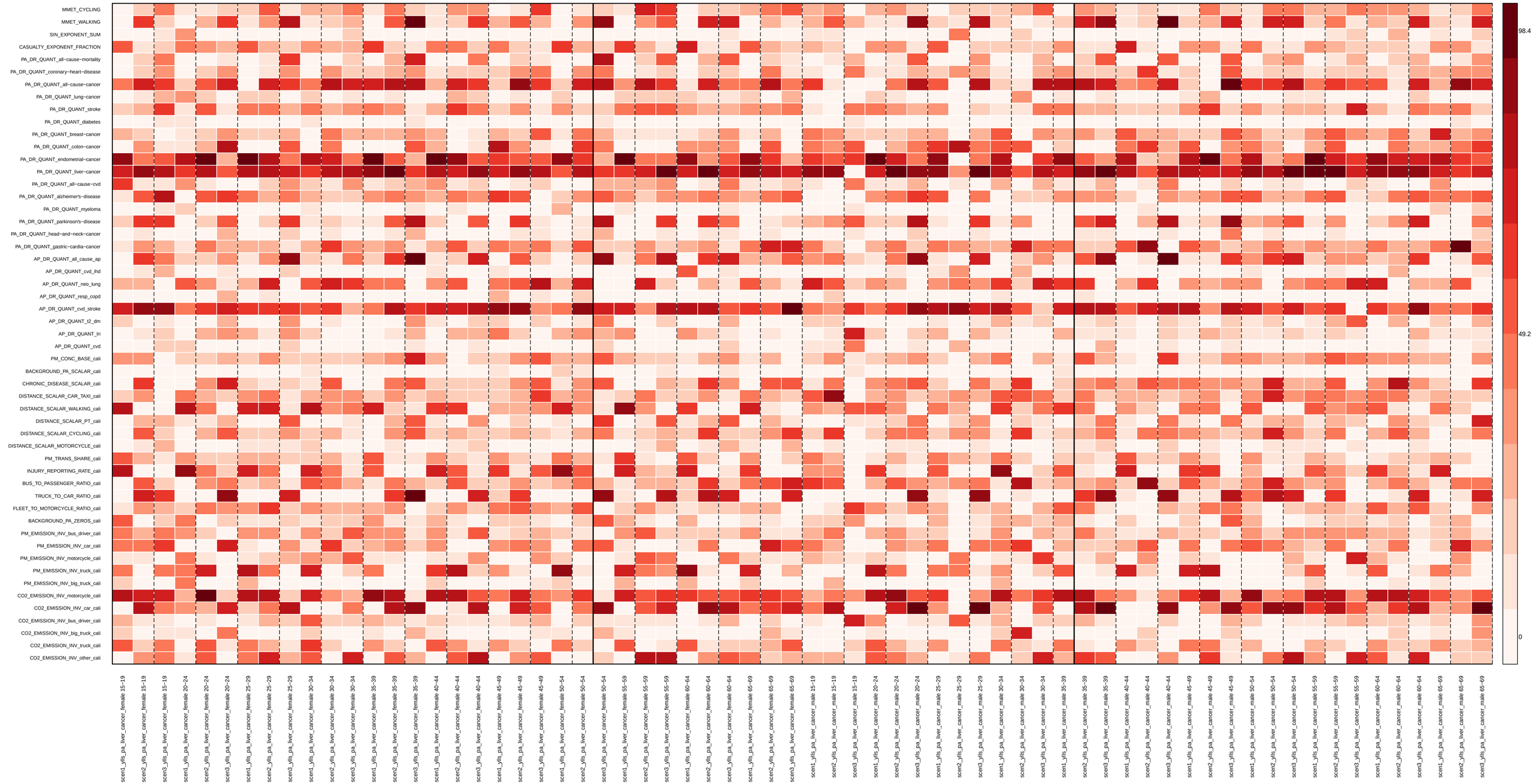


cali − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

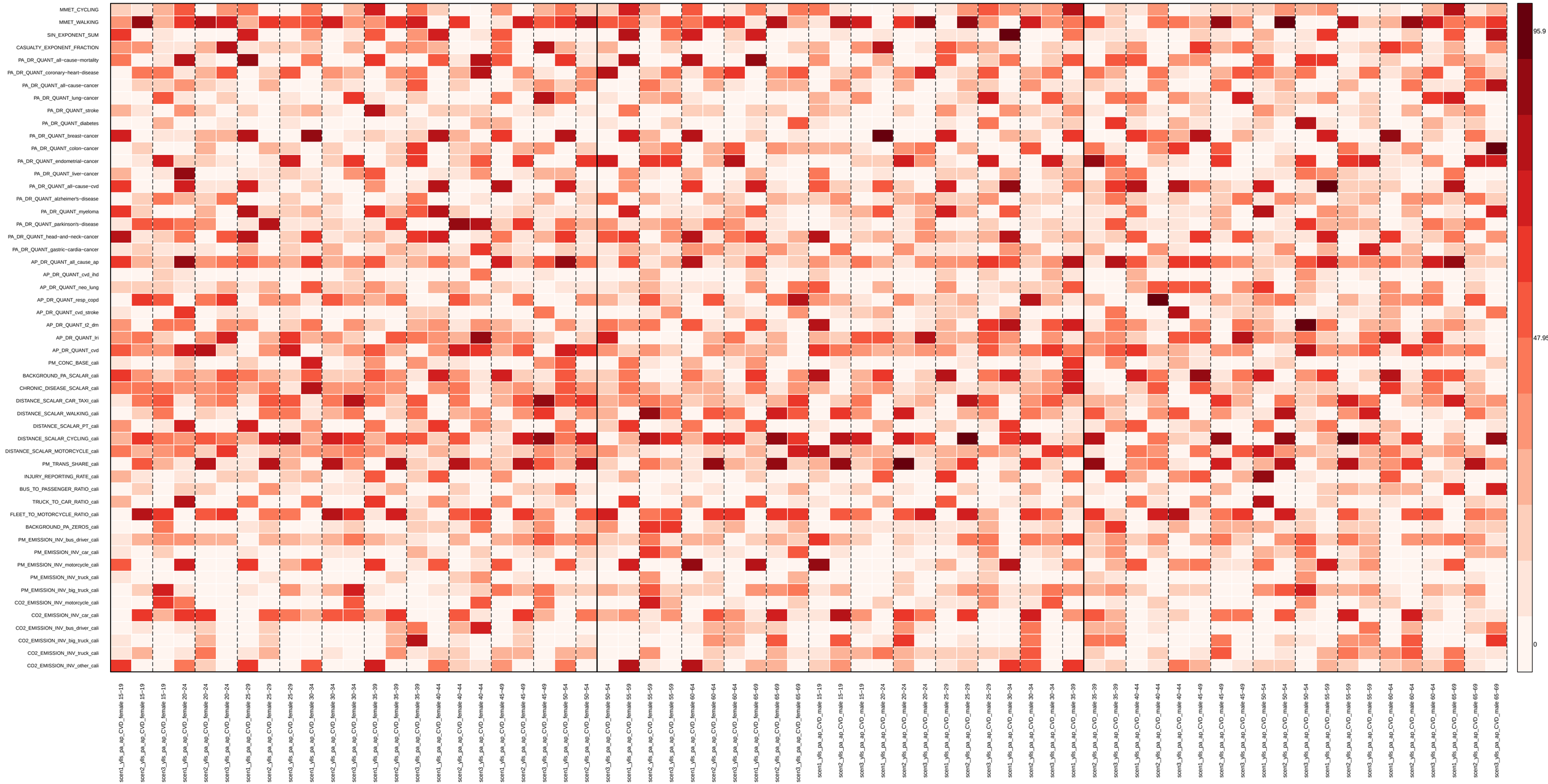


cali − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

cali − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

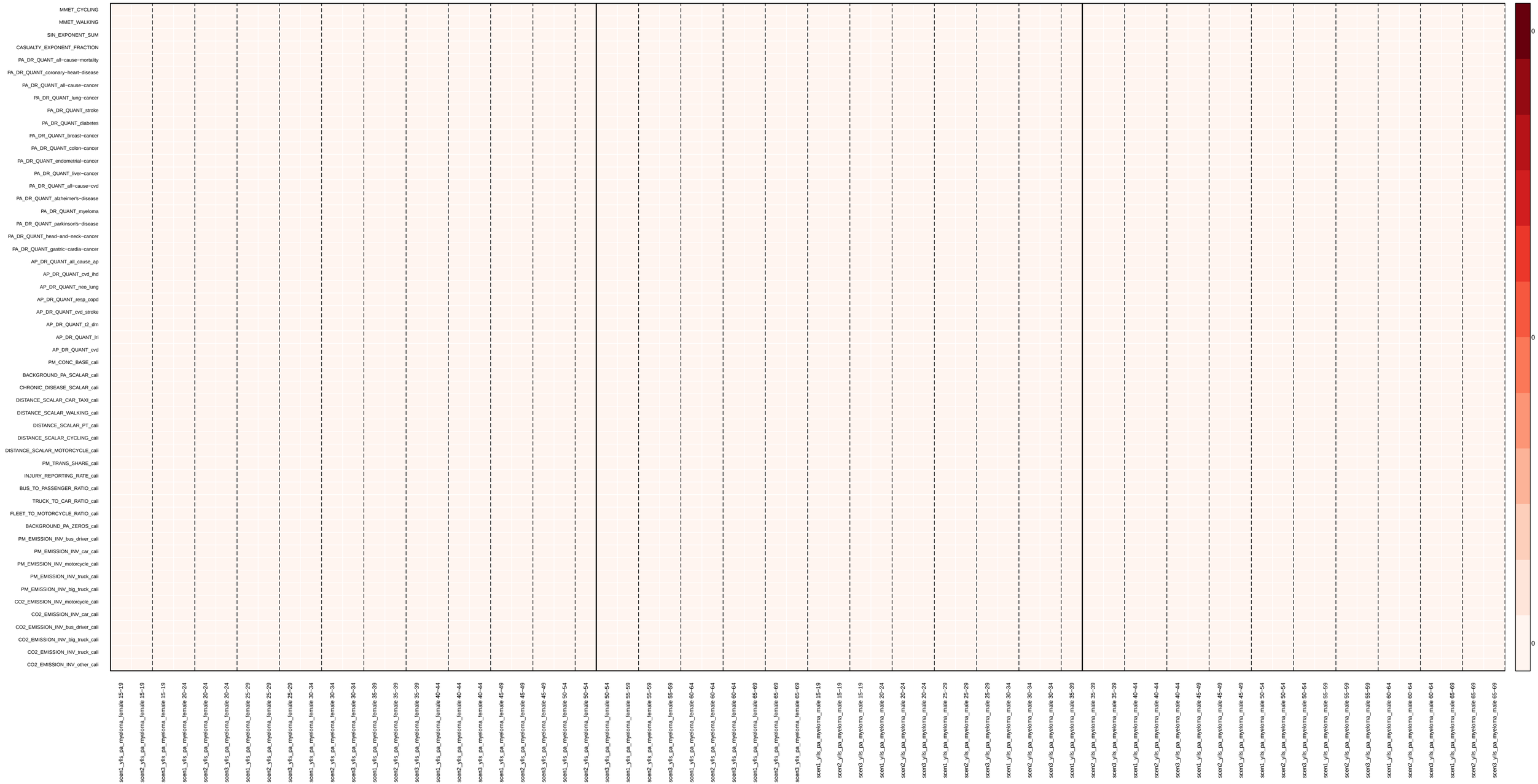


cali − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

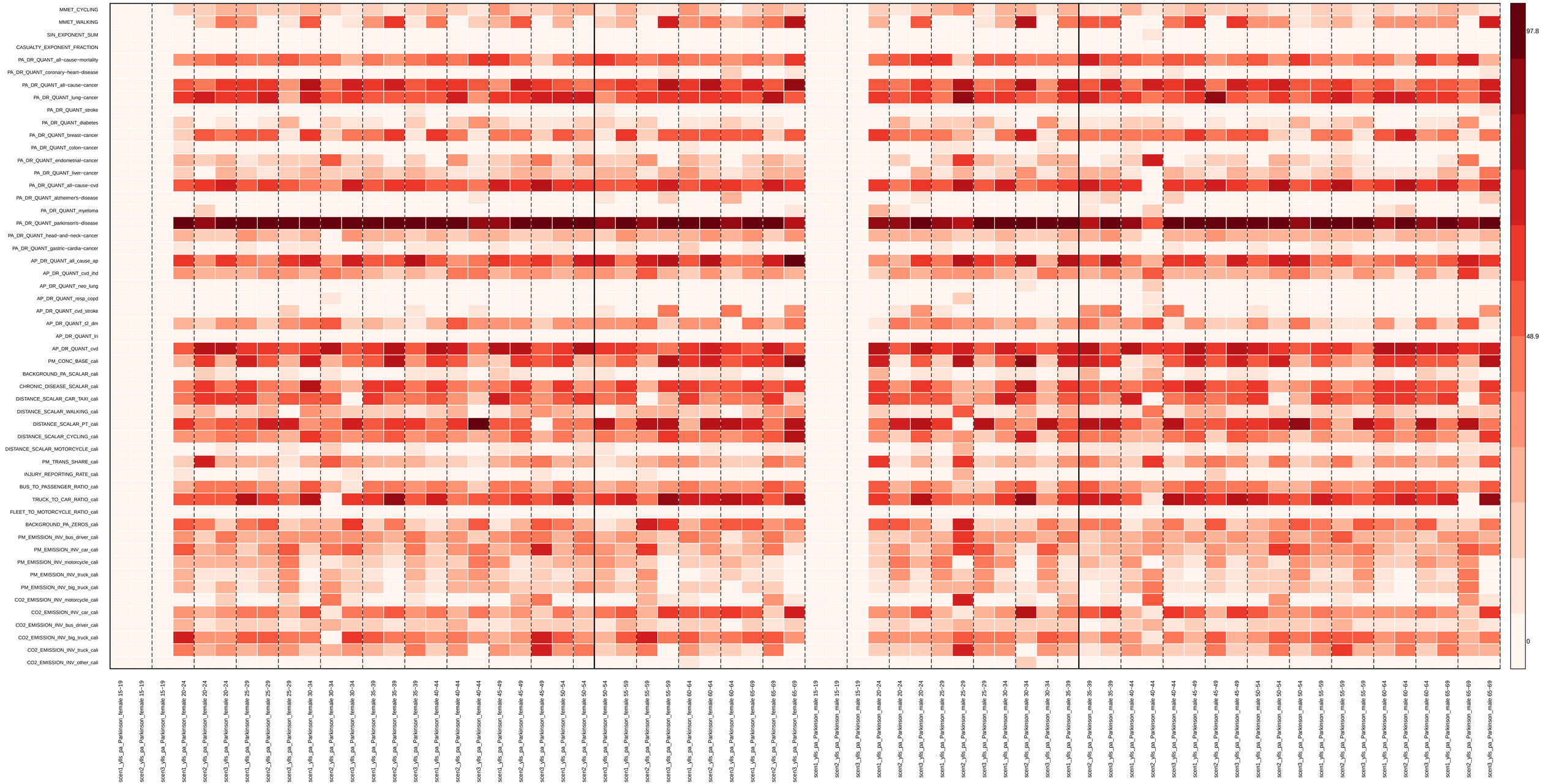


cali − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

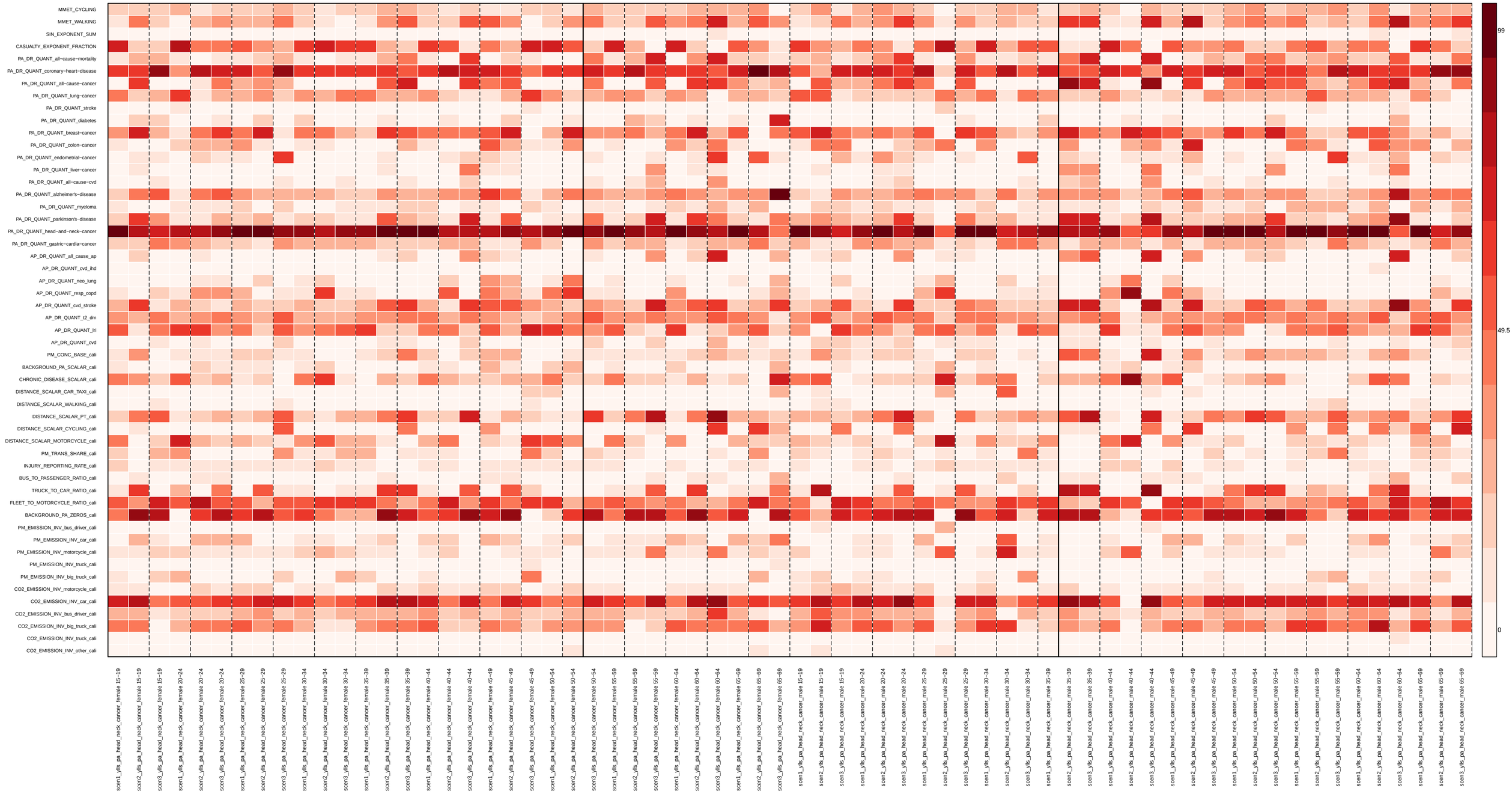
cali − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



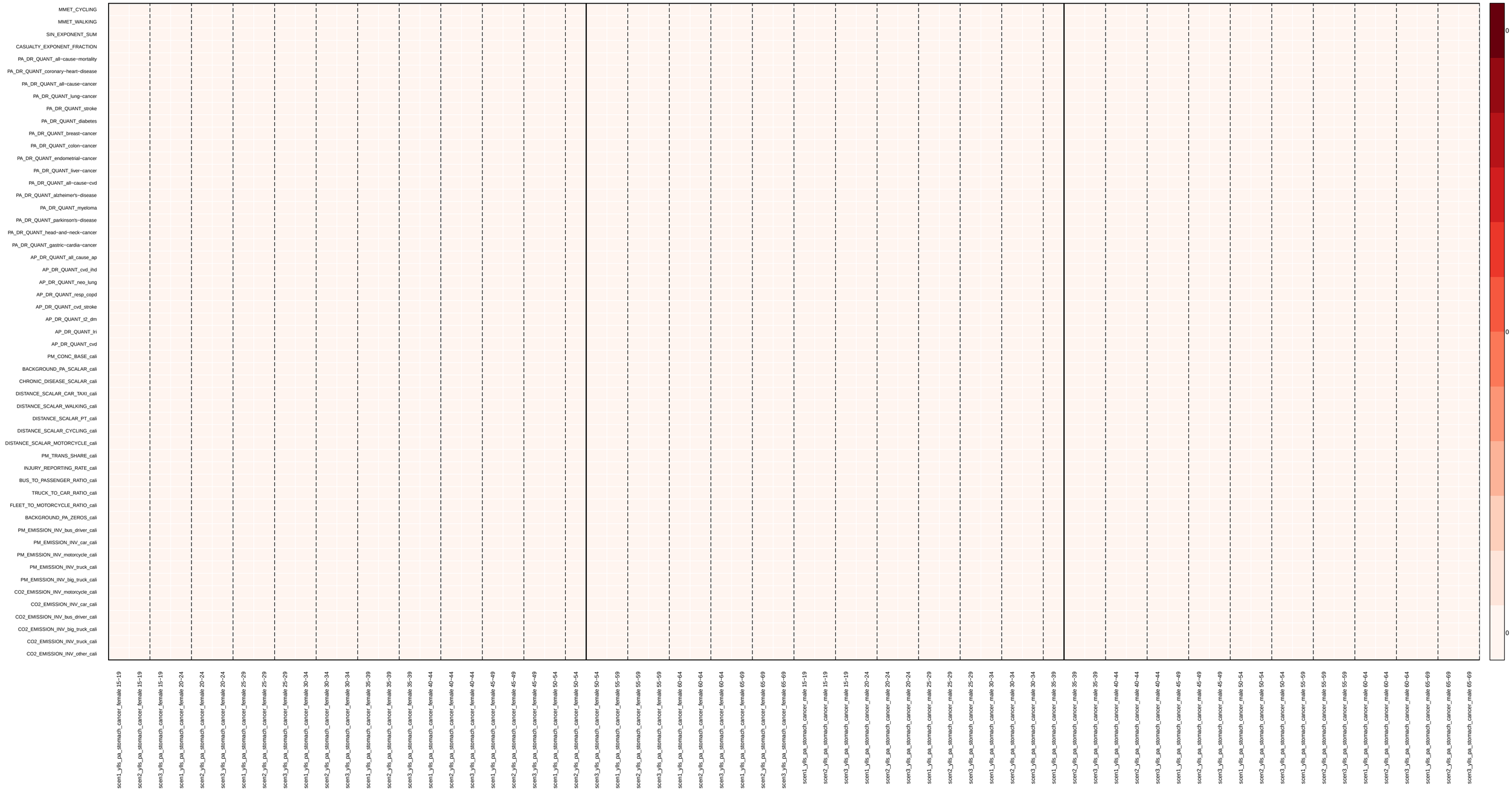
cali − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



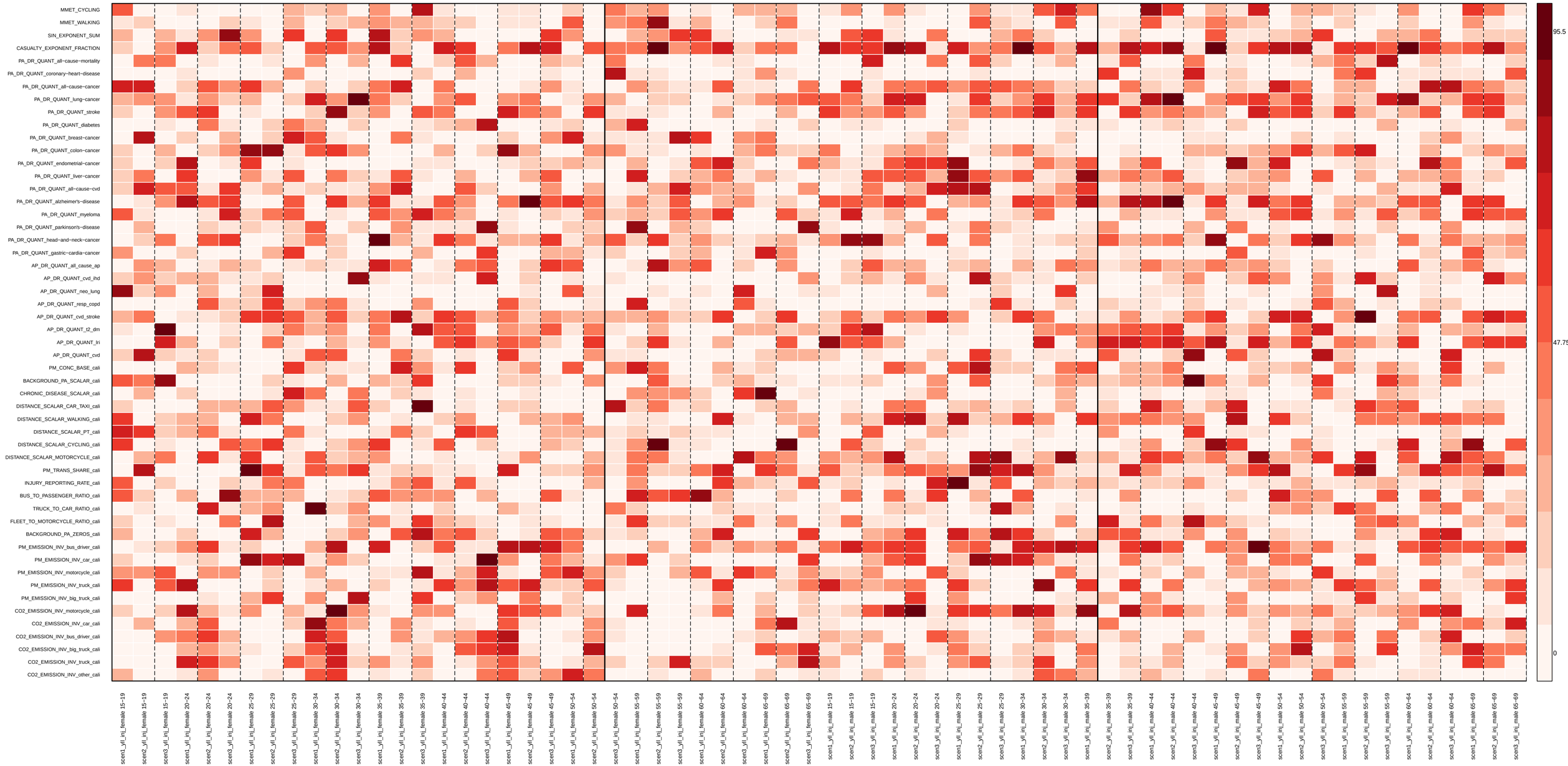
cali − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



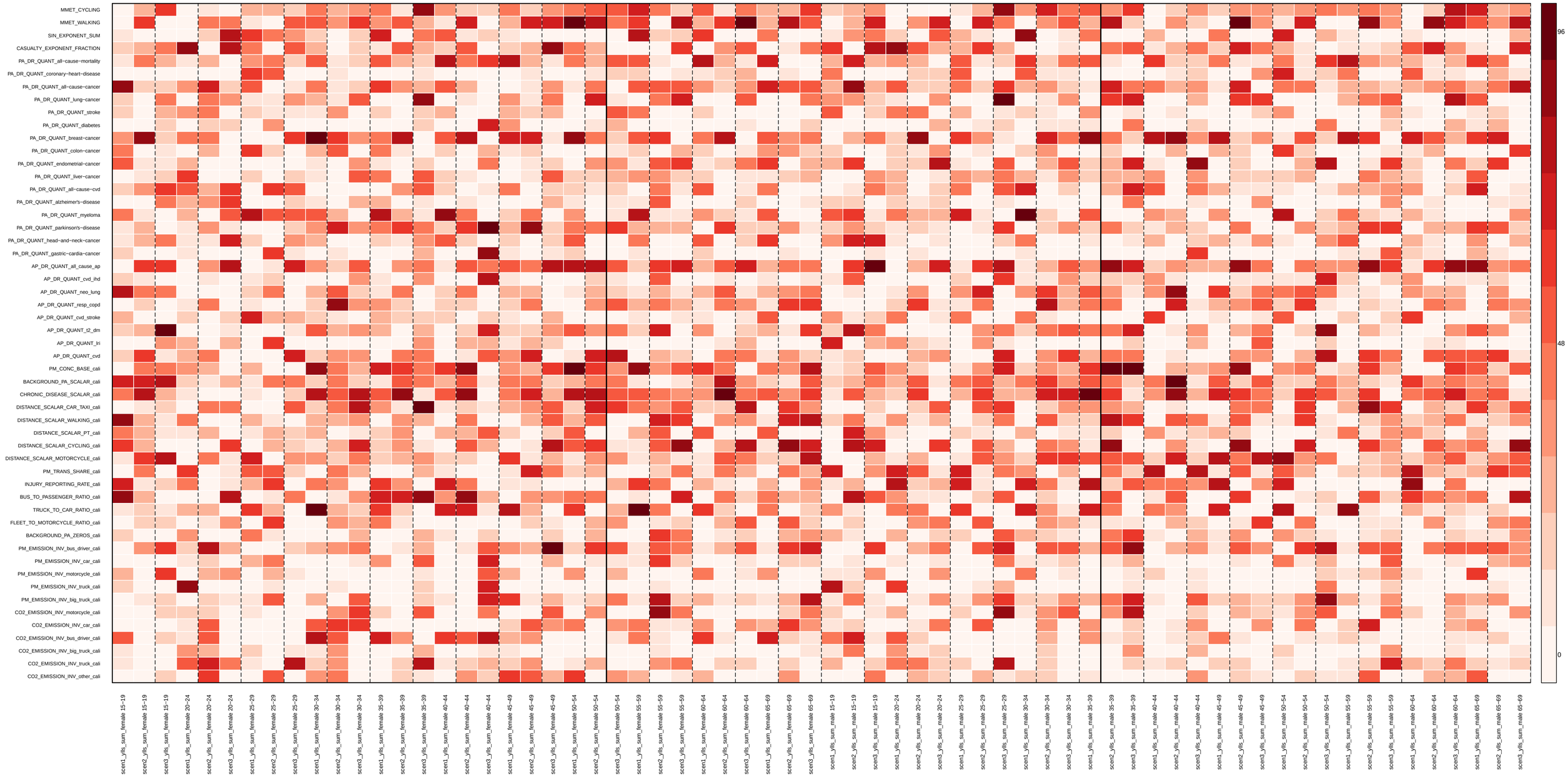
cali − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



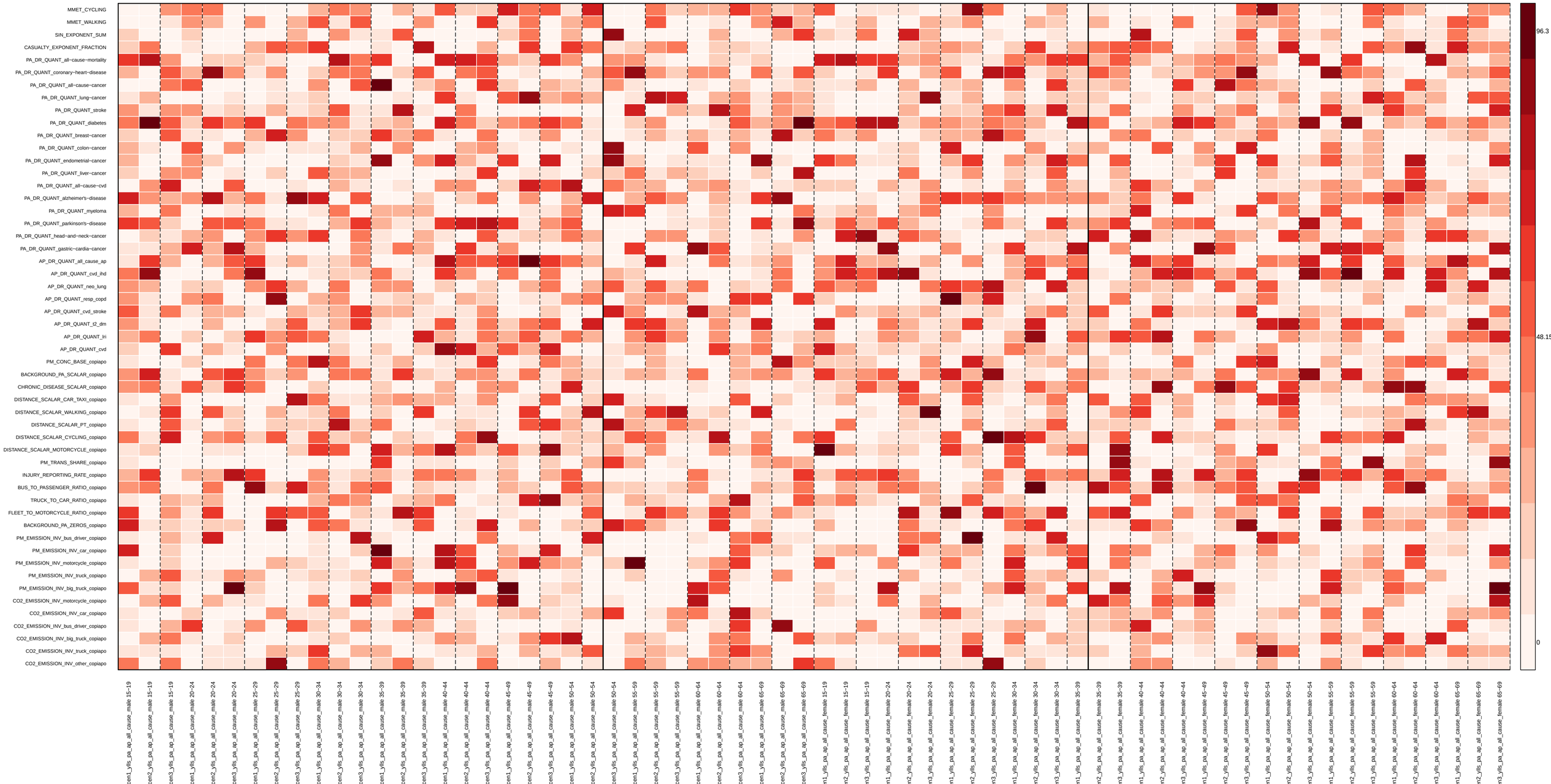
cali − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



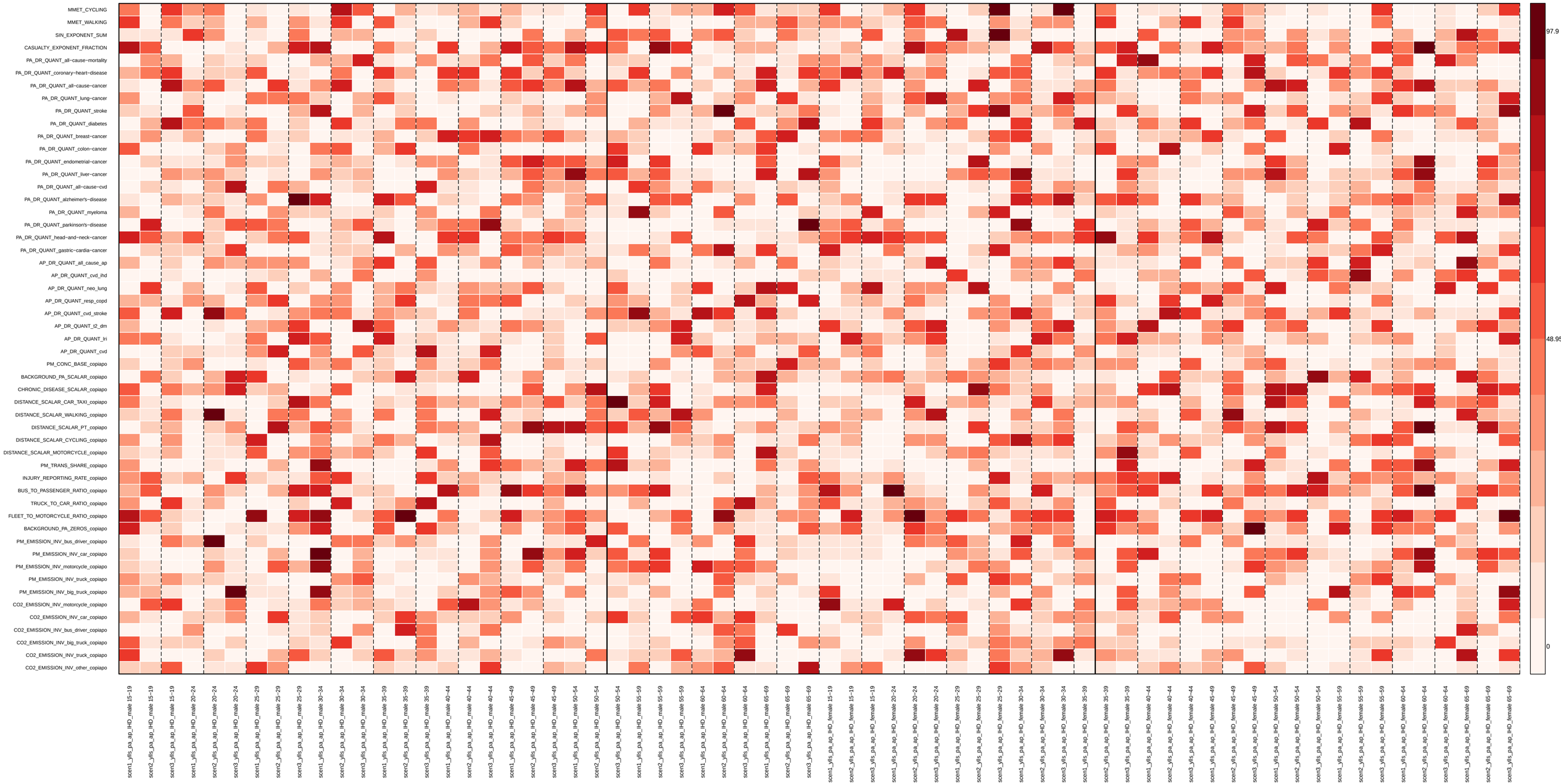
cali − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



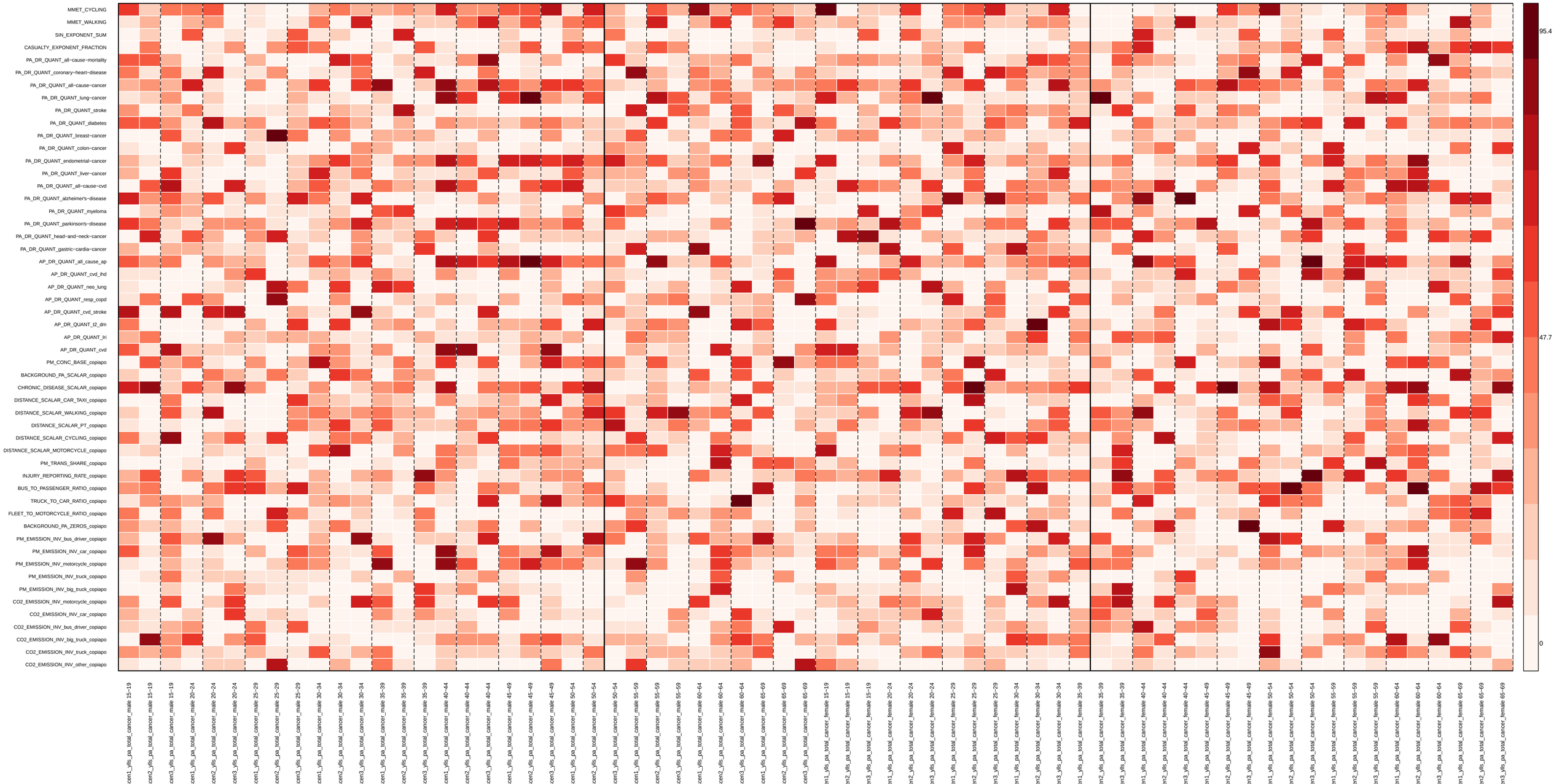
copiapo − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



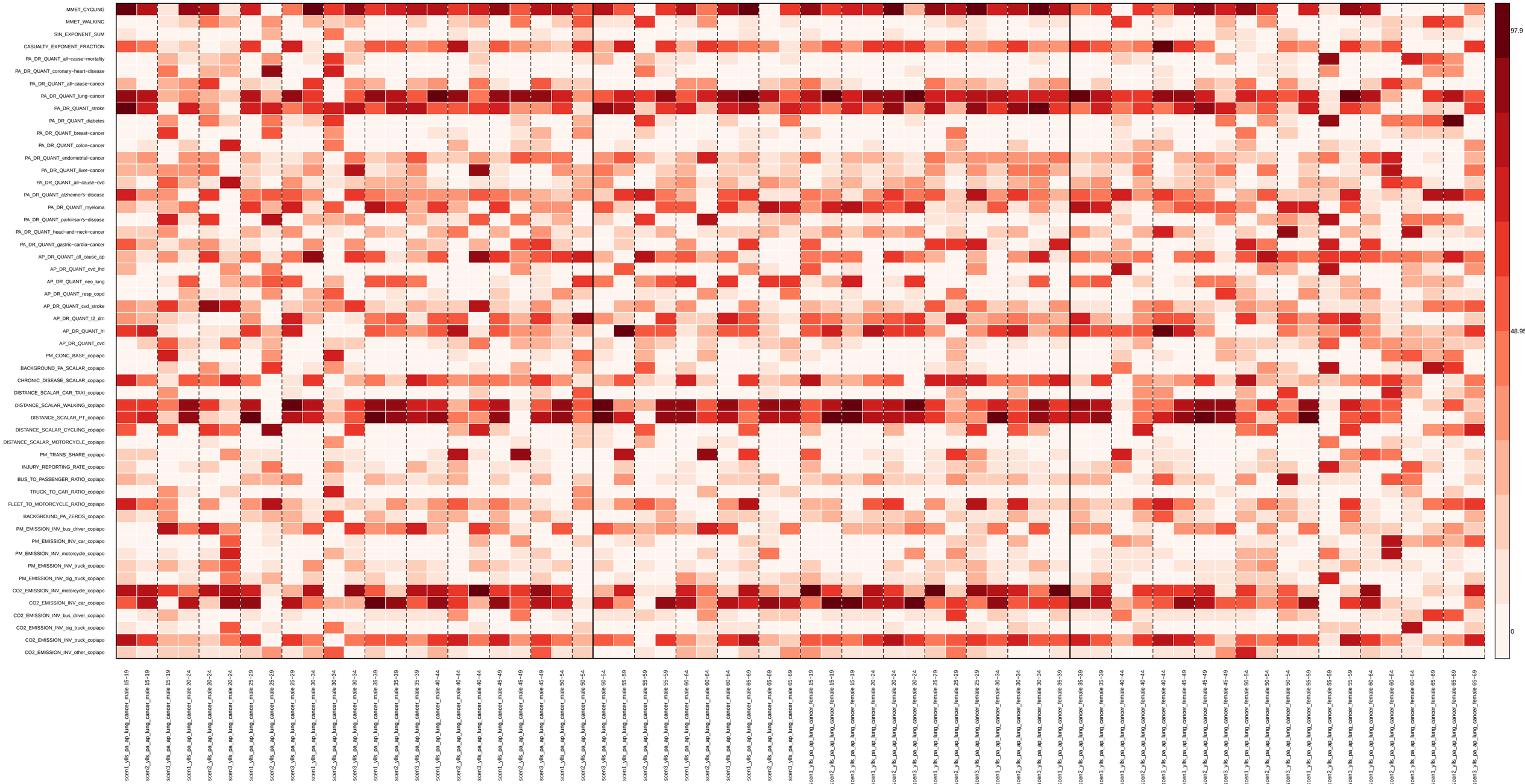
copiapo − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



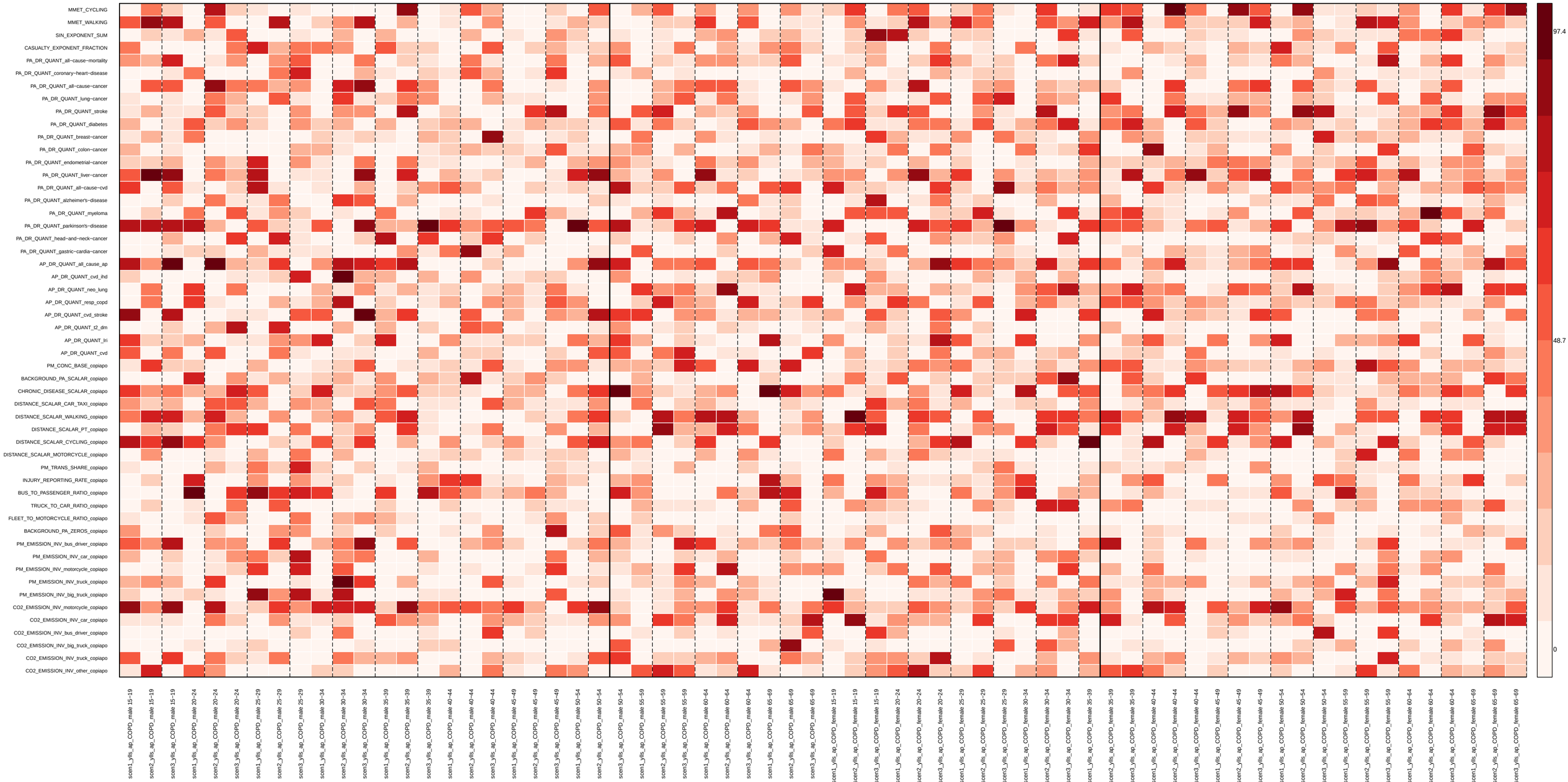
copiapo − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



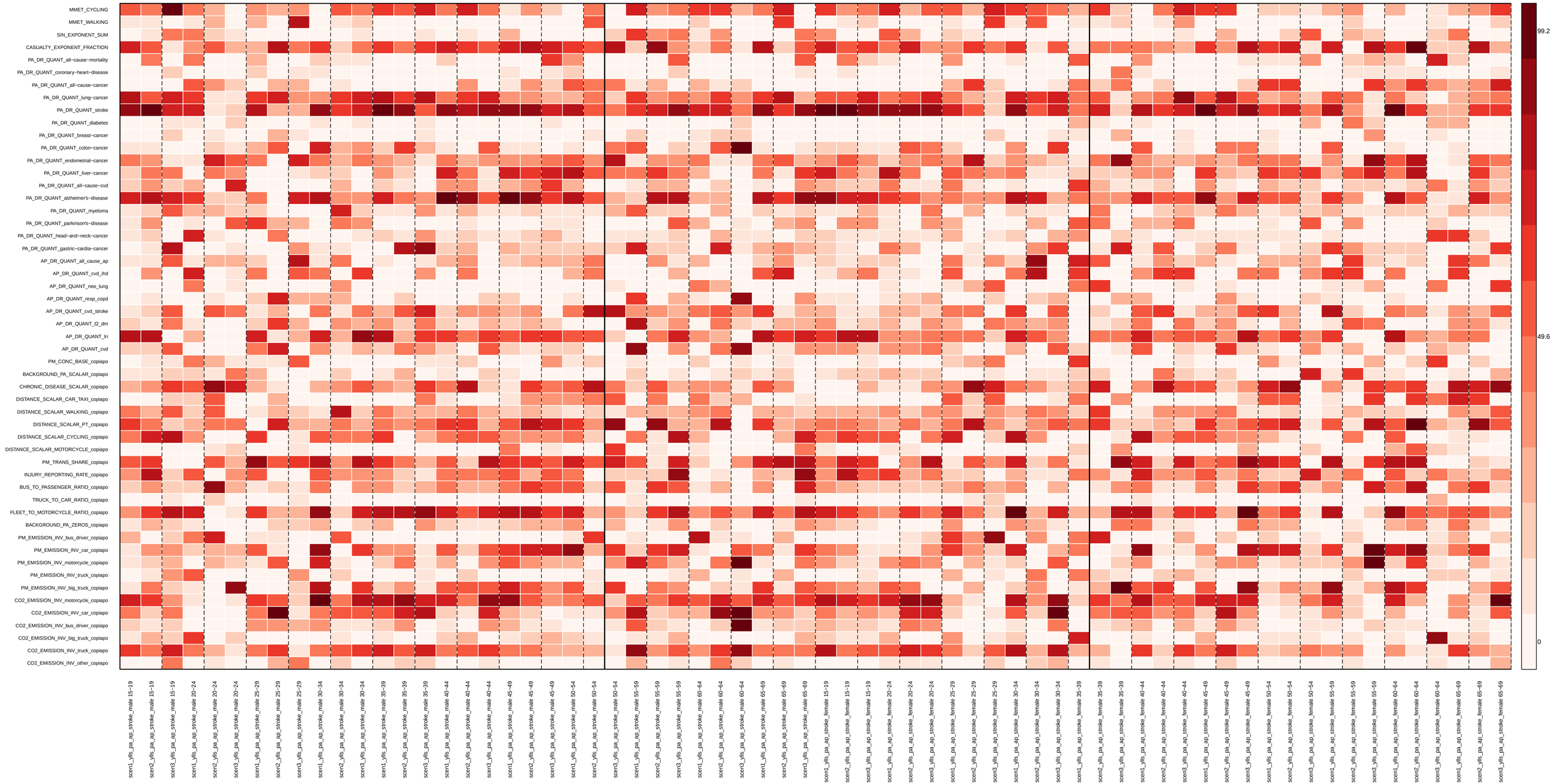
copiapo − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



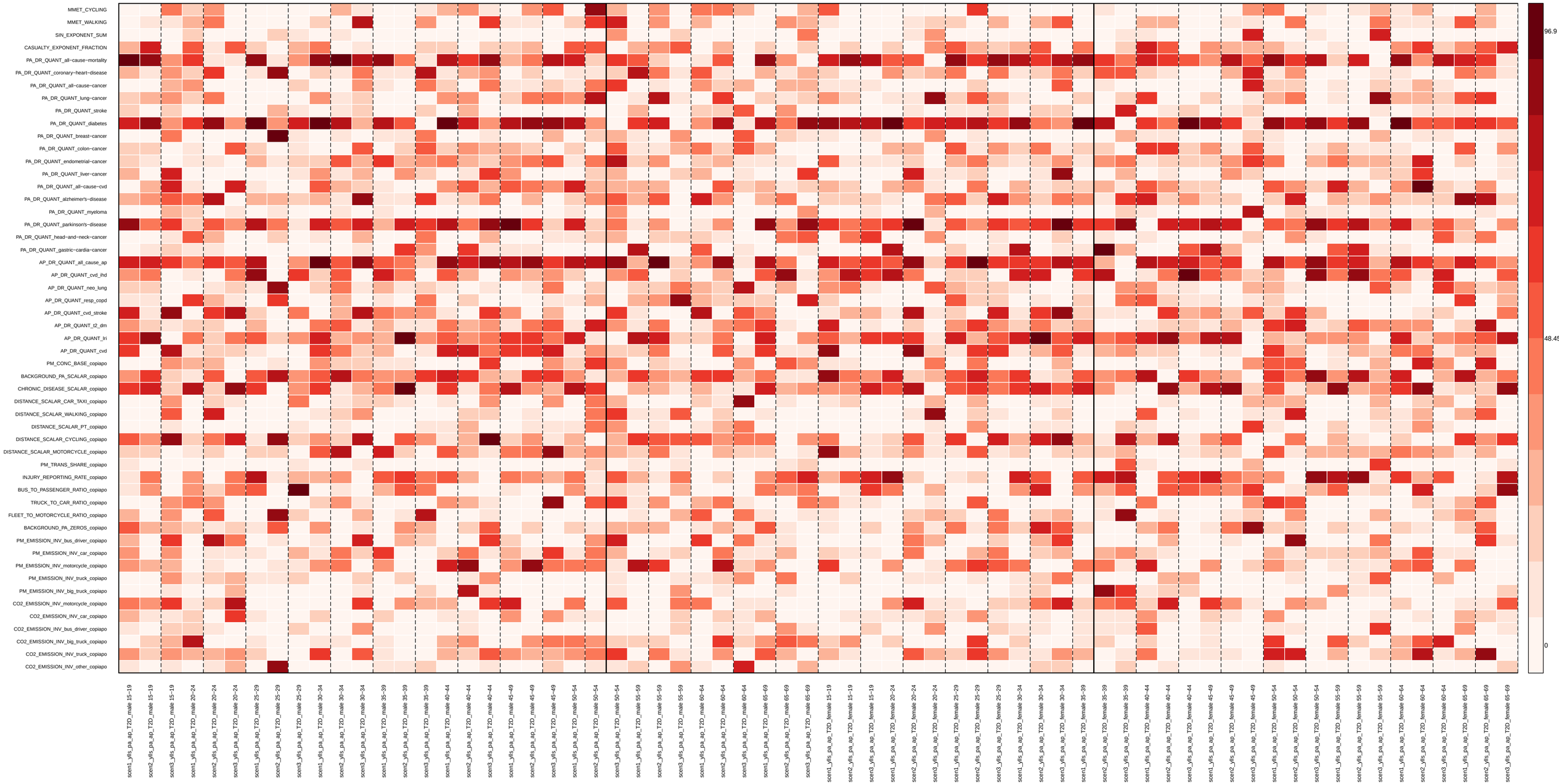
copiapo − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



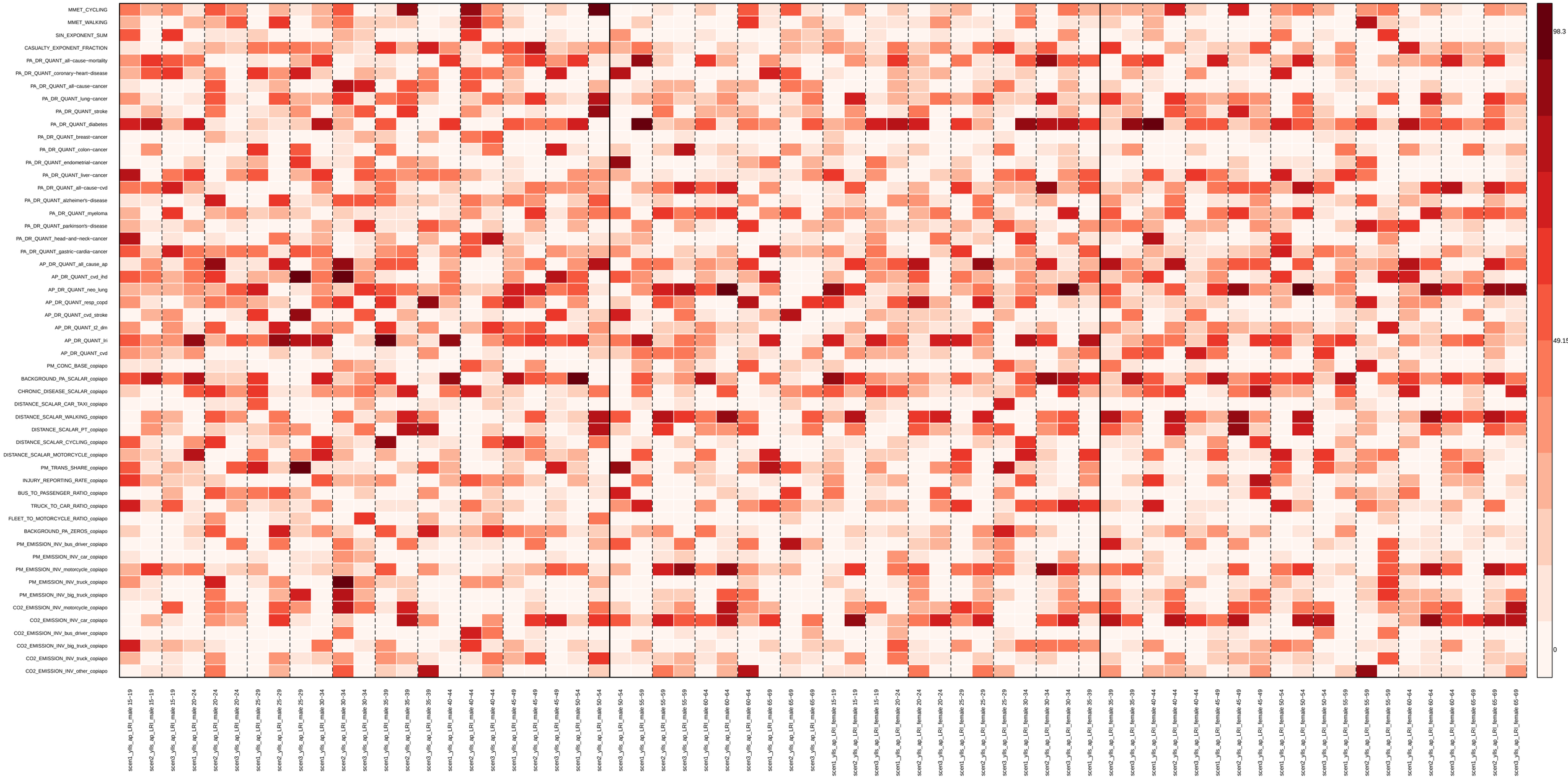
copiapo − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



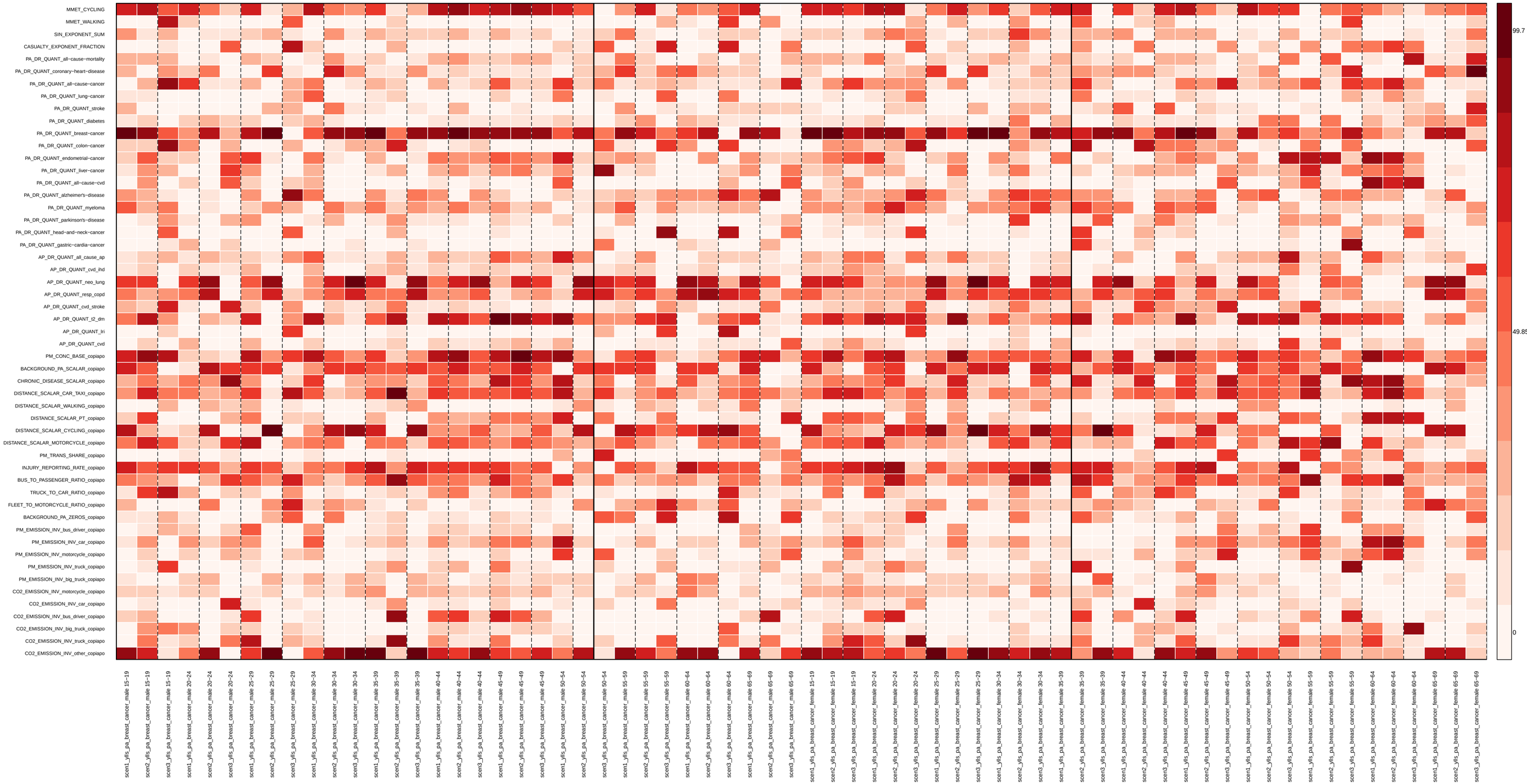
copiapo − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



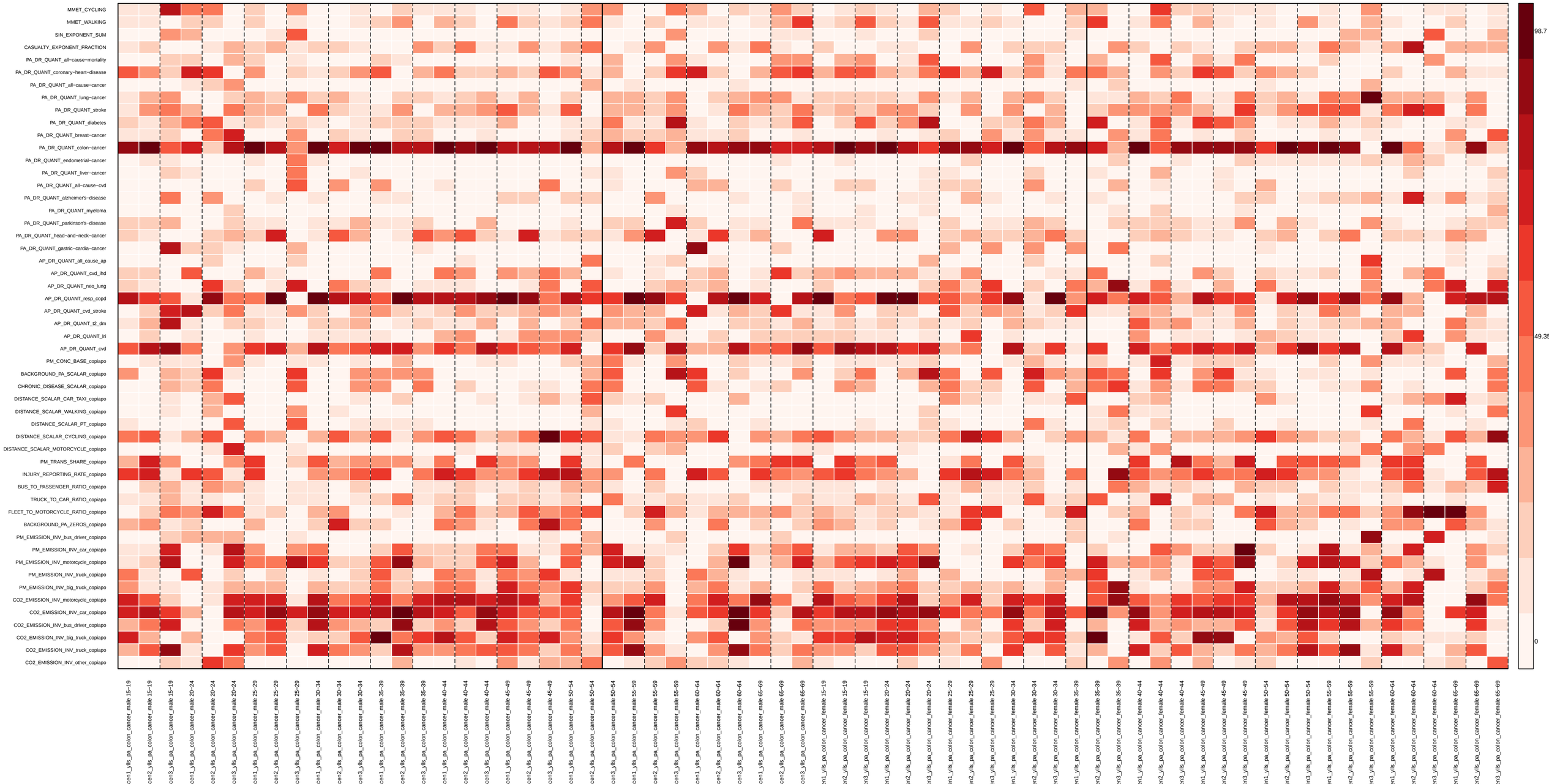
copiapo − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



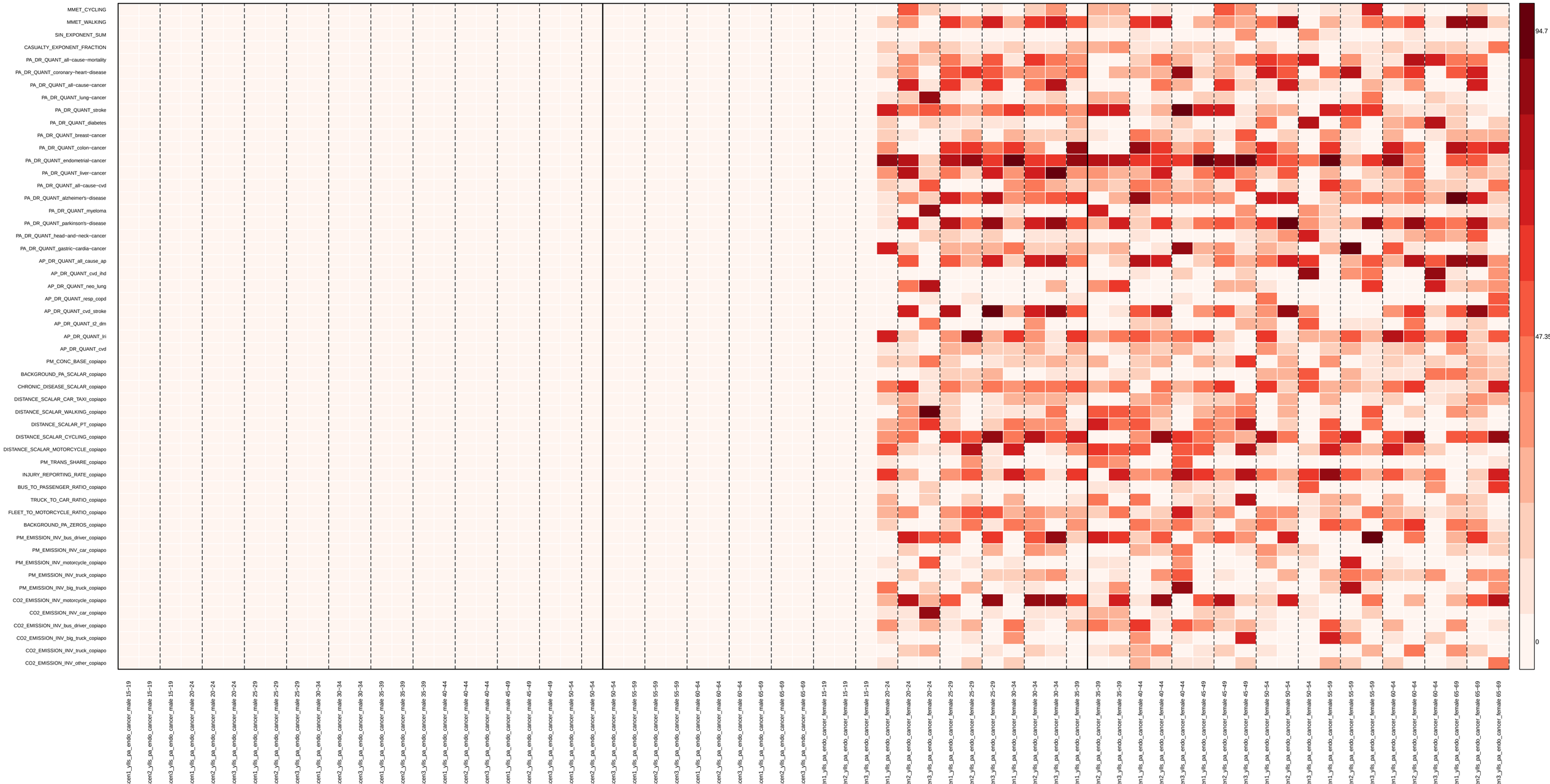
copiapo − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



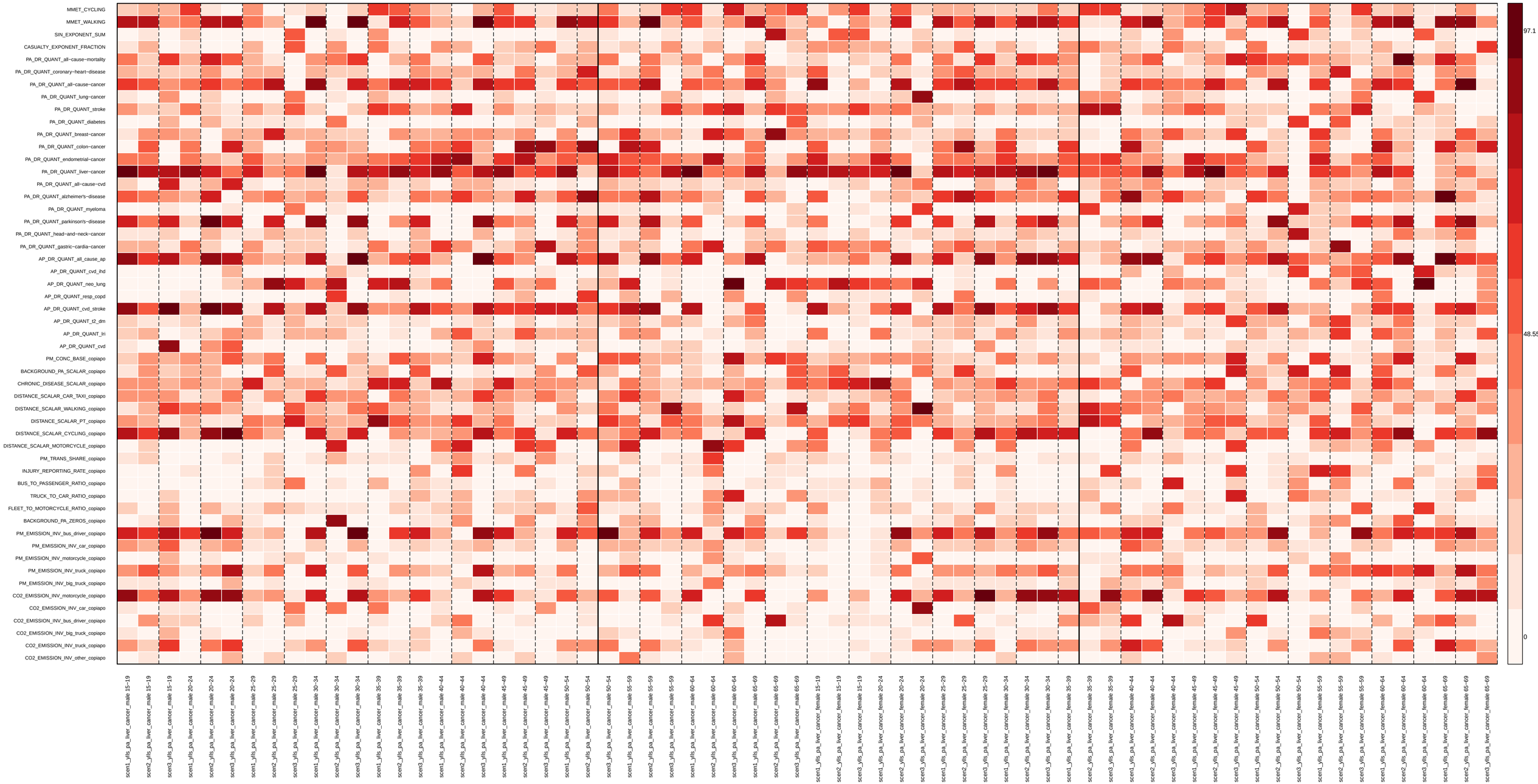
copiapo − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



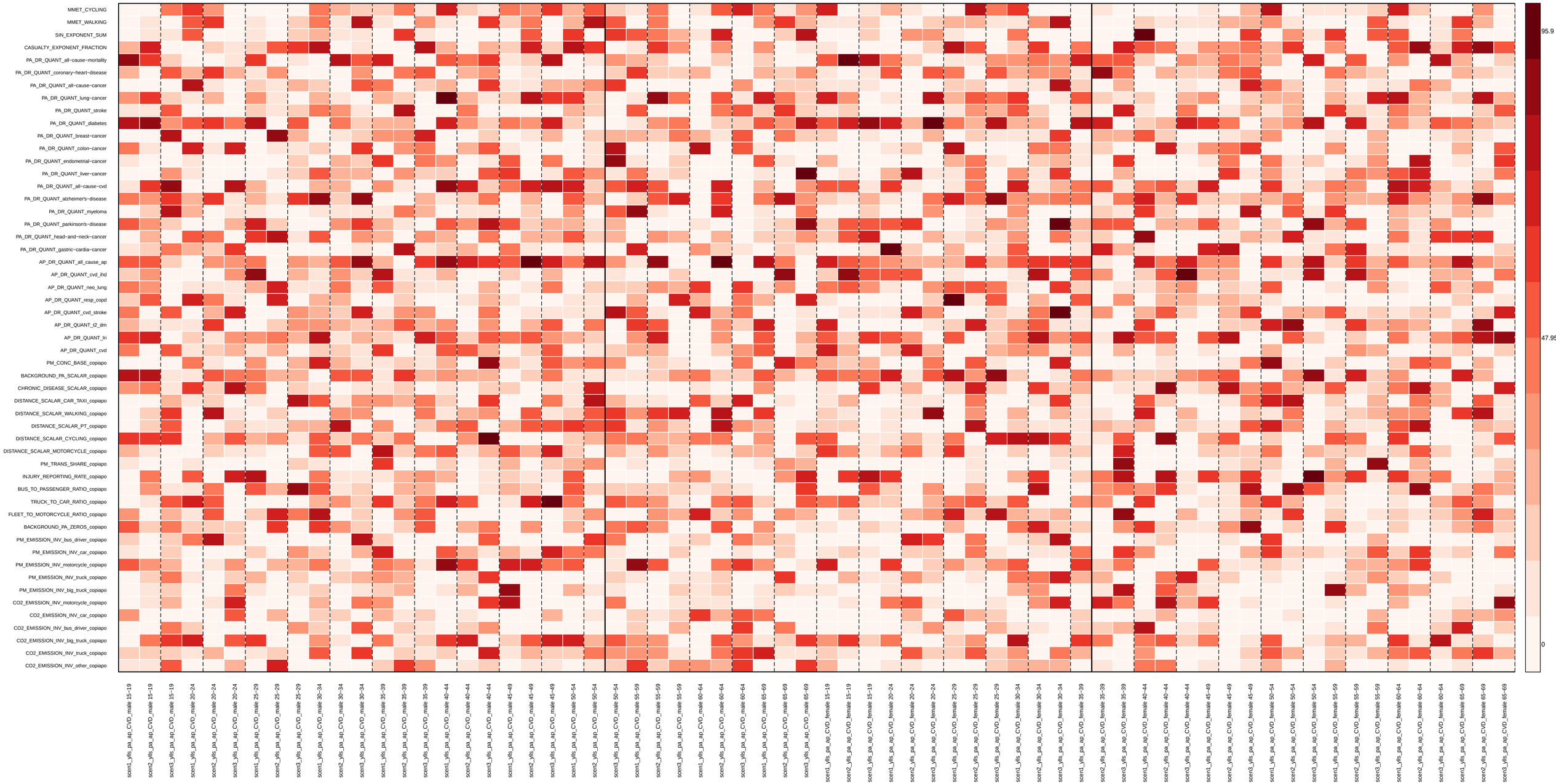
copiapo − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



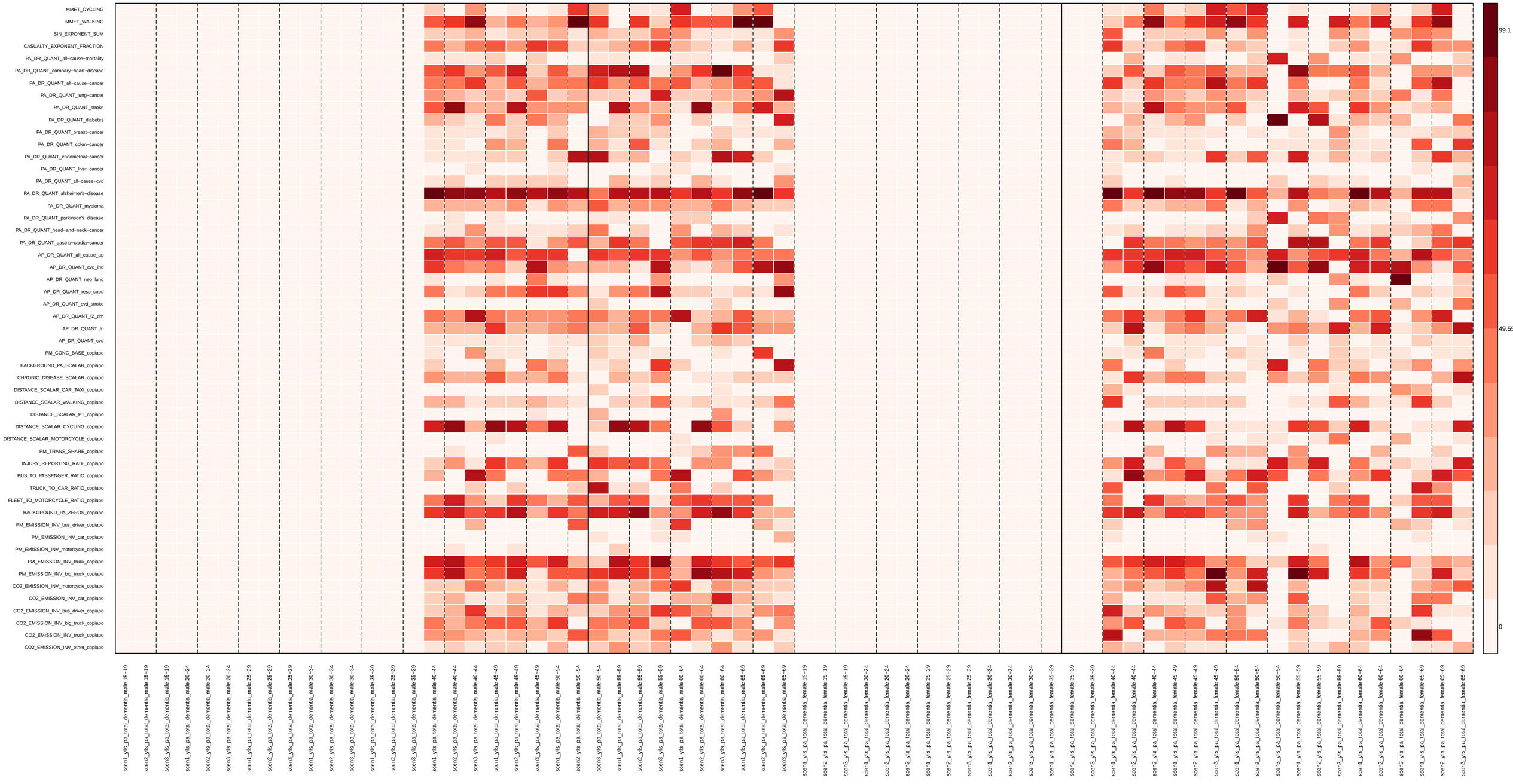
copiapo − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



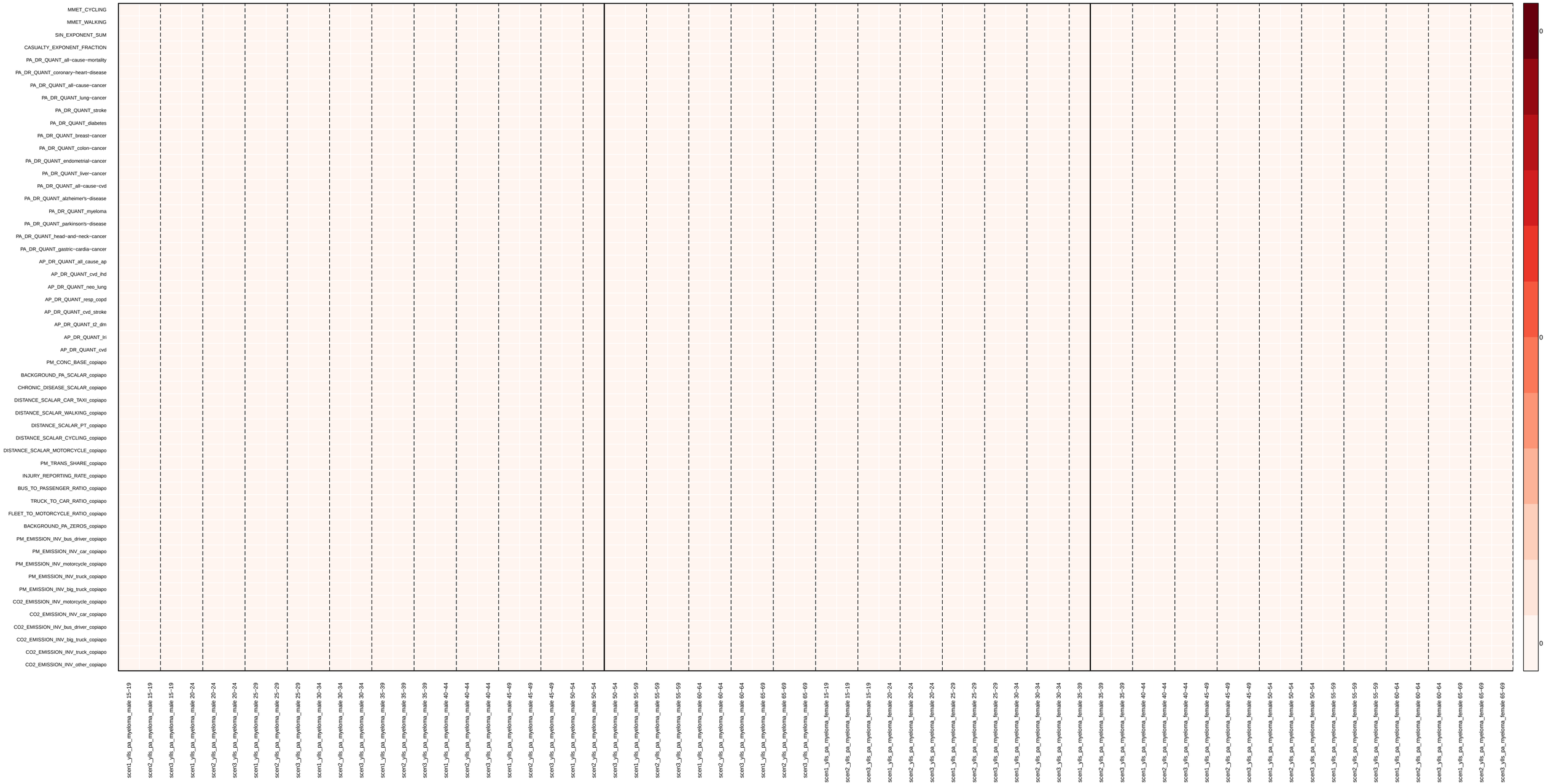
copiapo − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



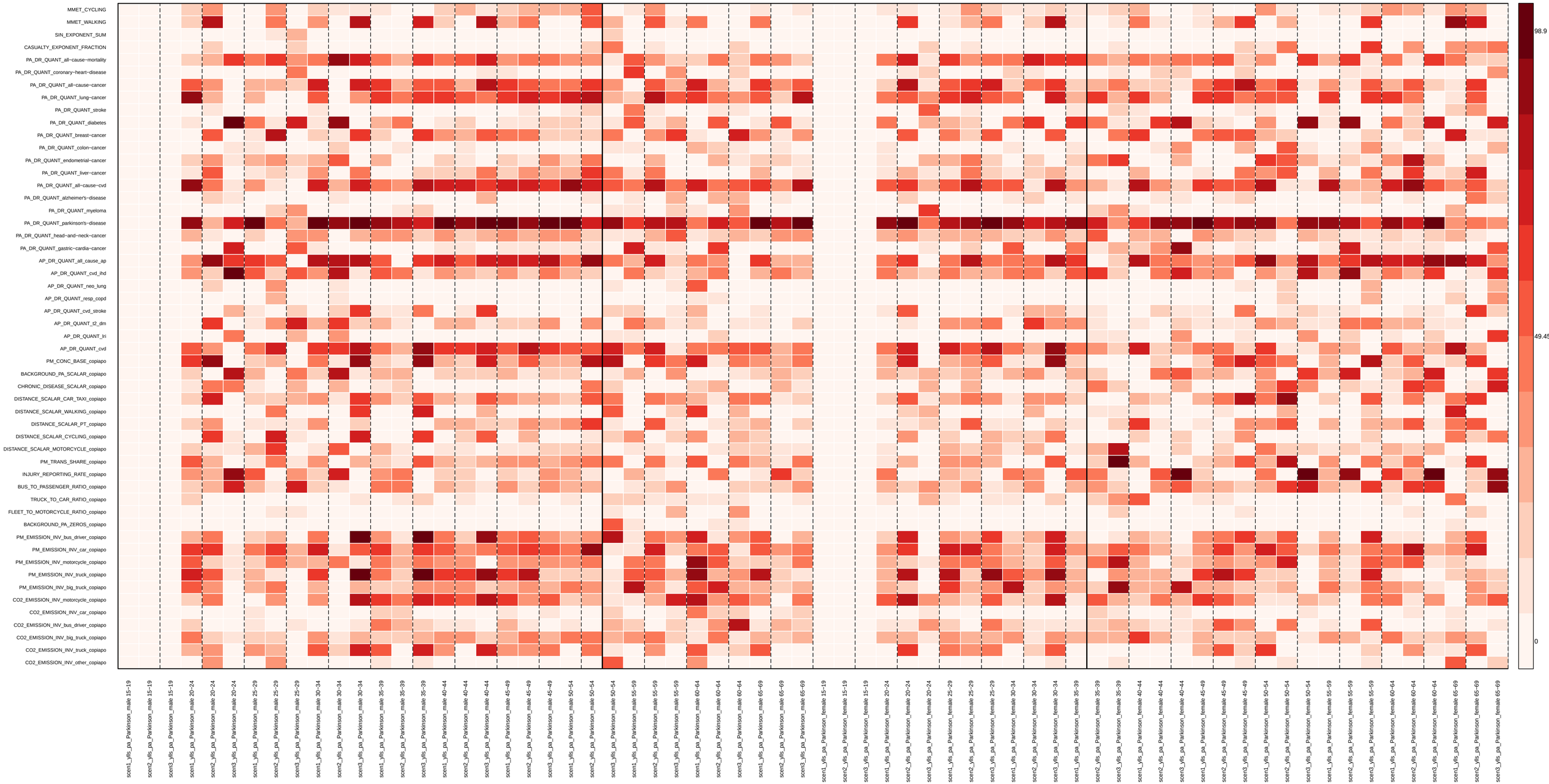
copiapo − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



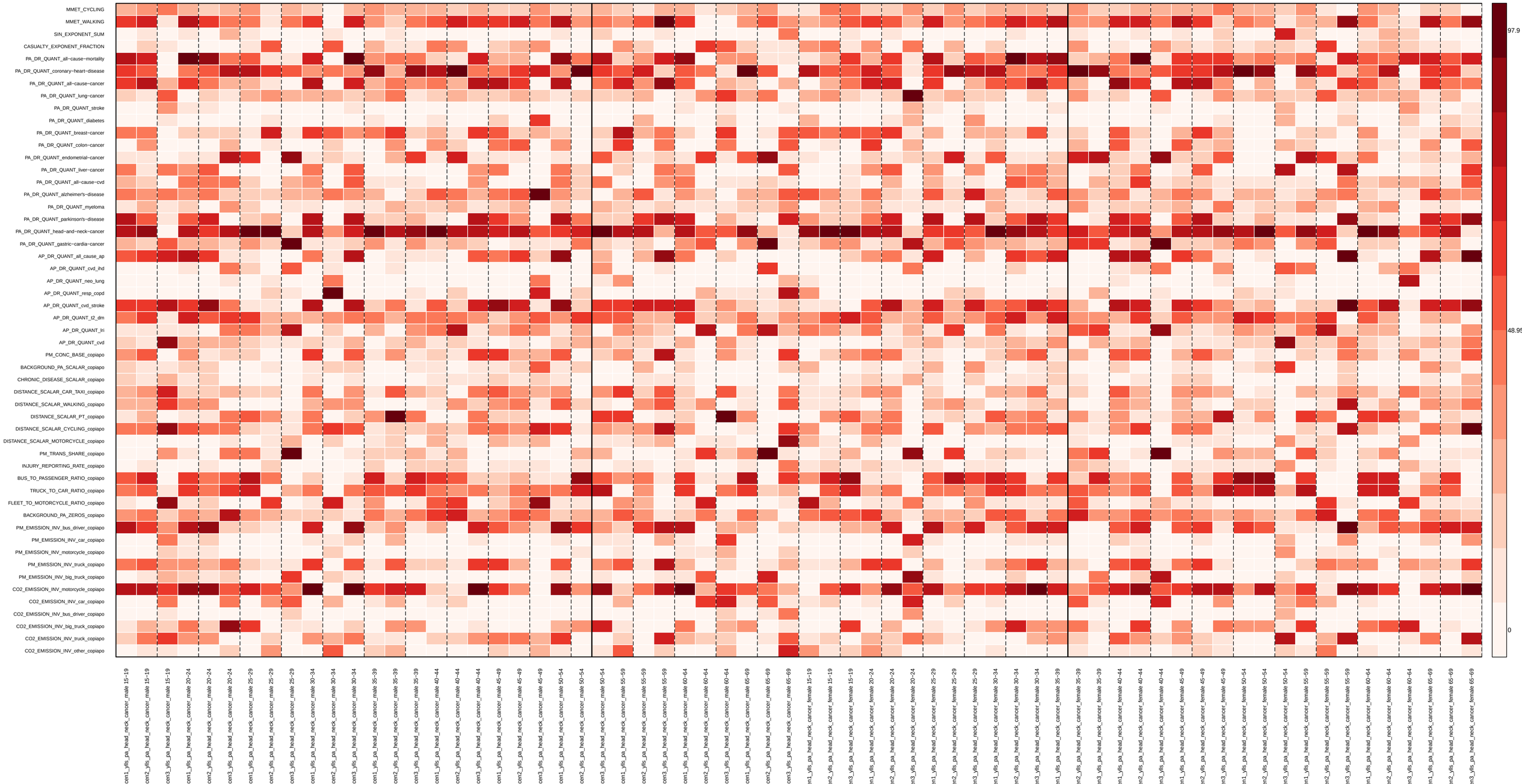
copiapo − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



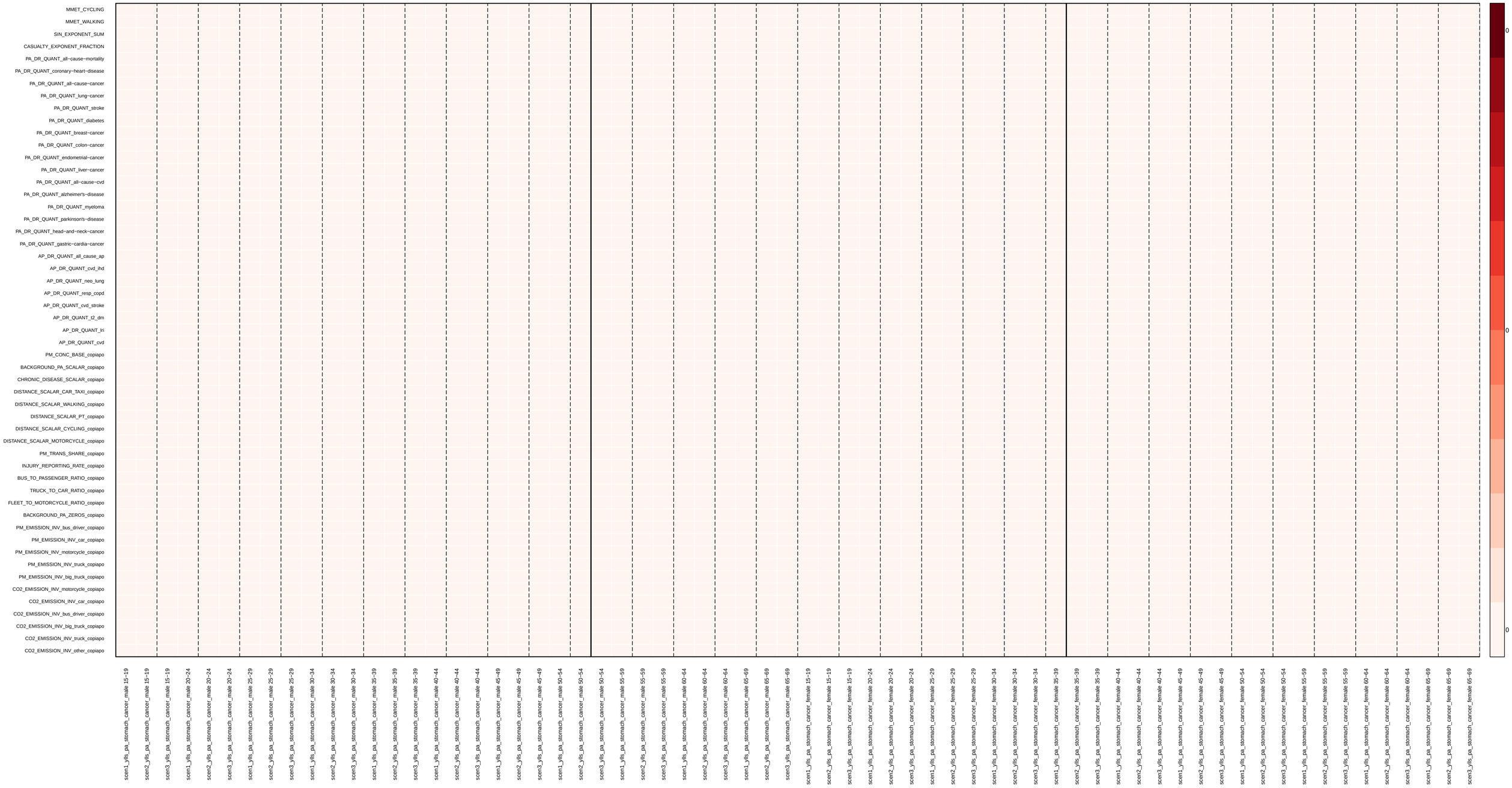
copiapo − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



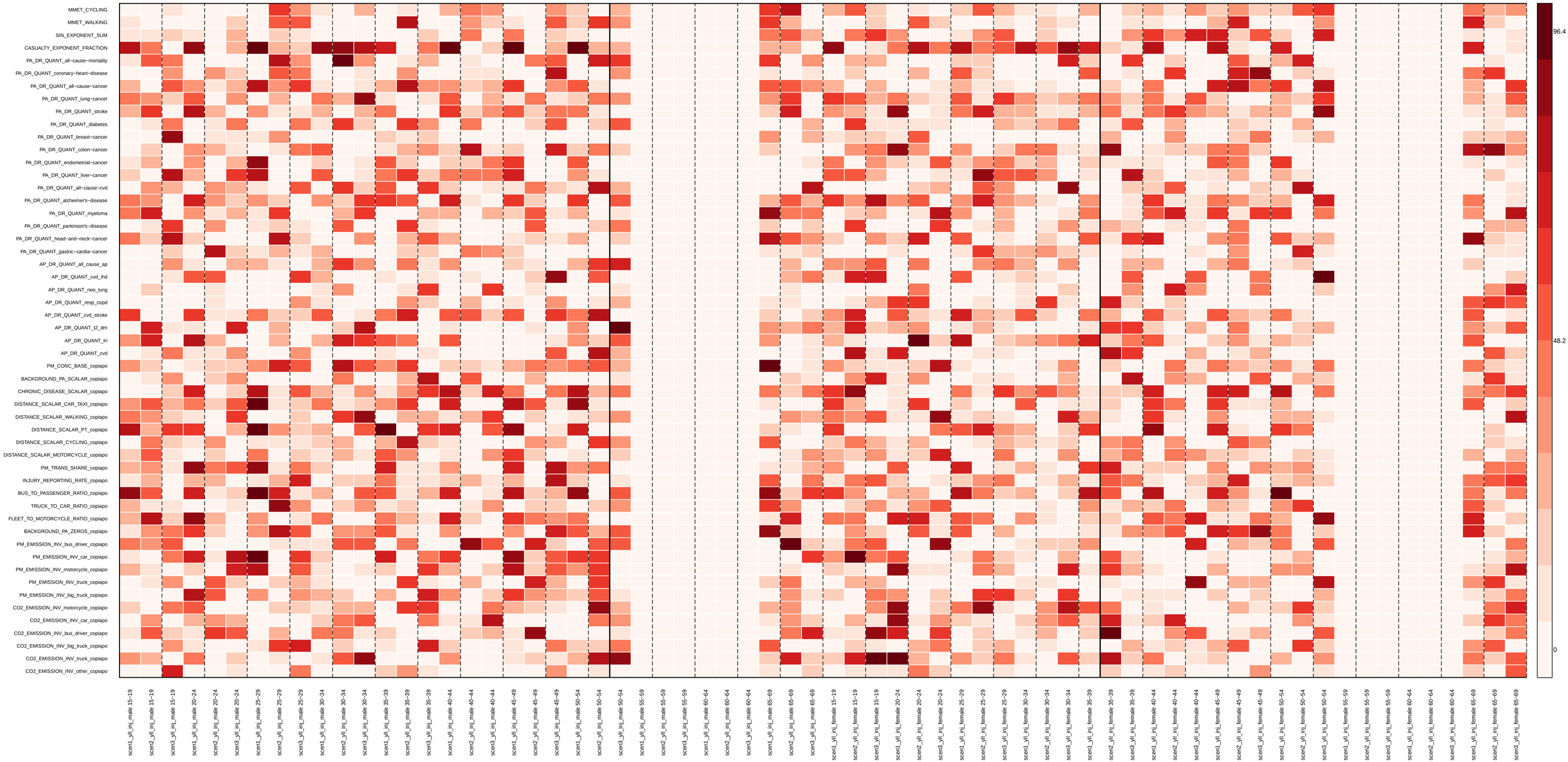
copiapo − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



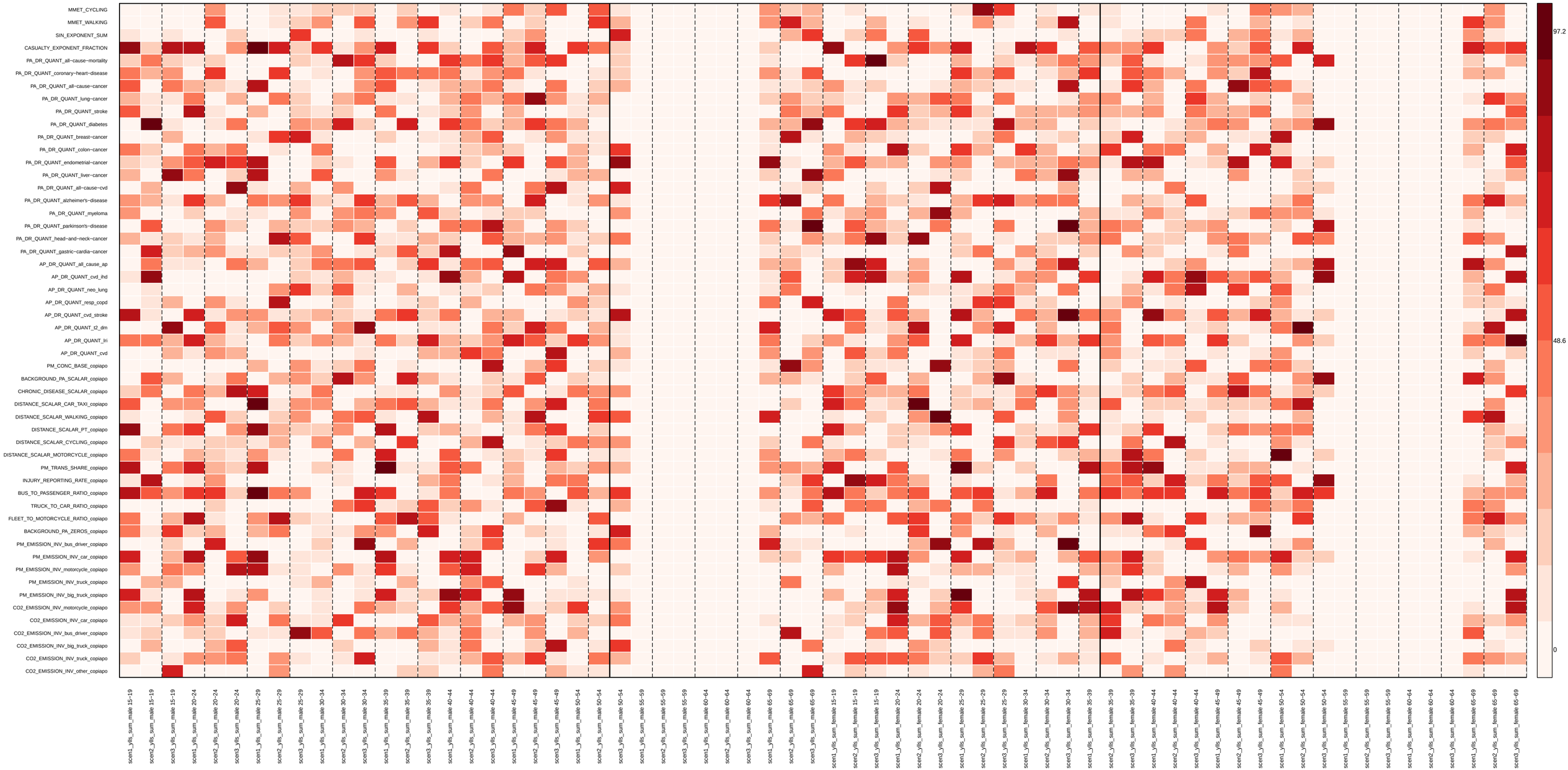
copiapo − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



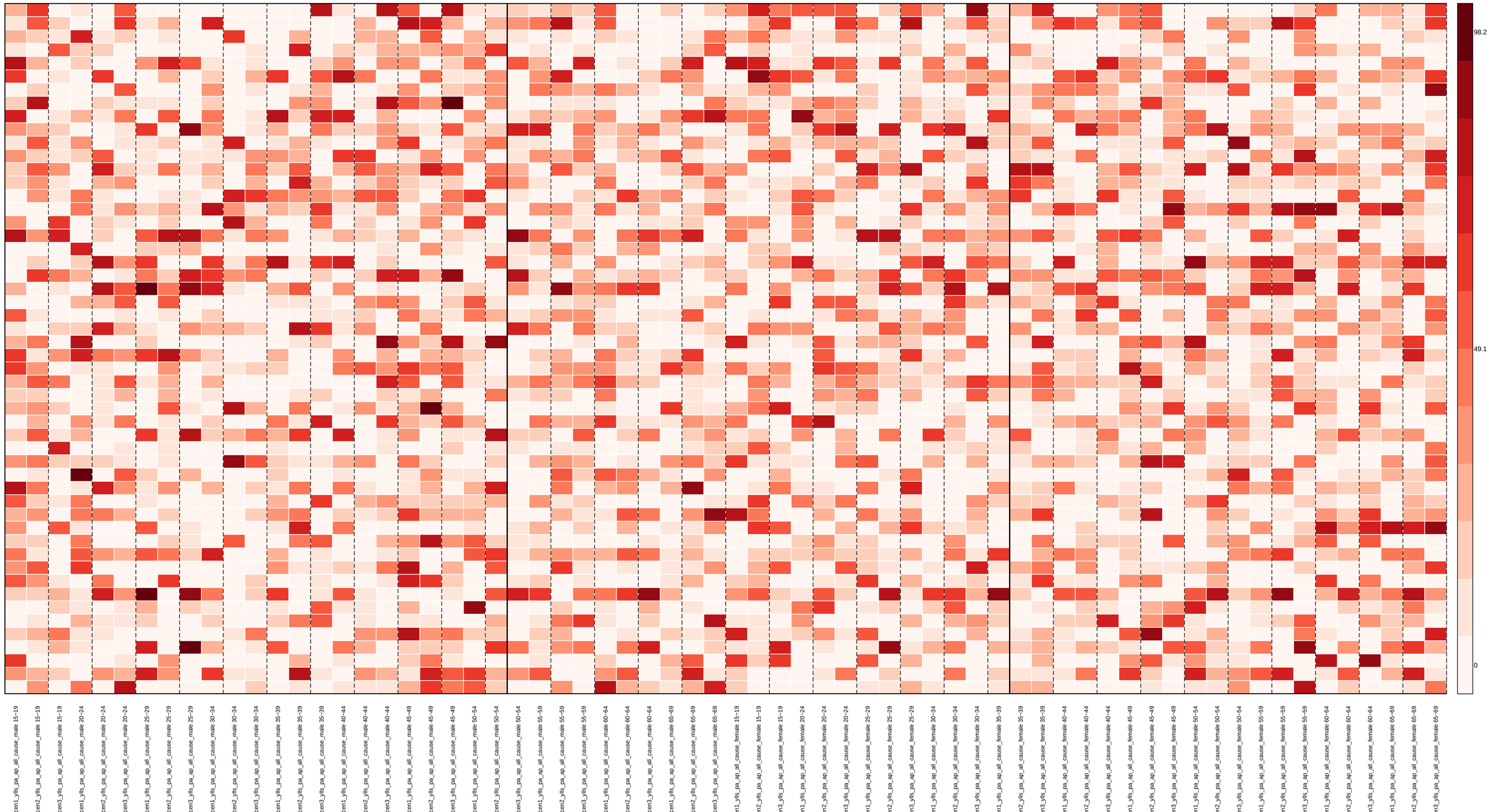
copiapo − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



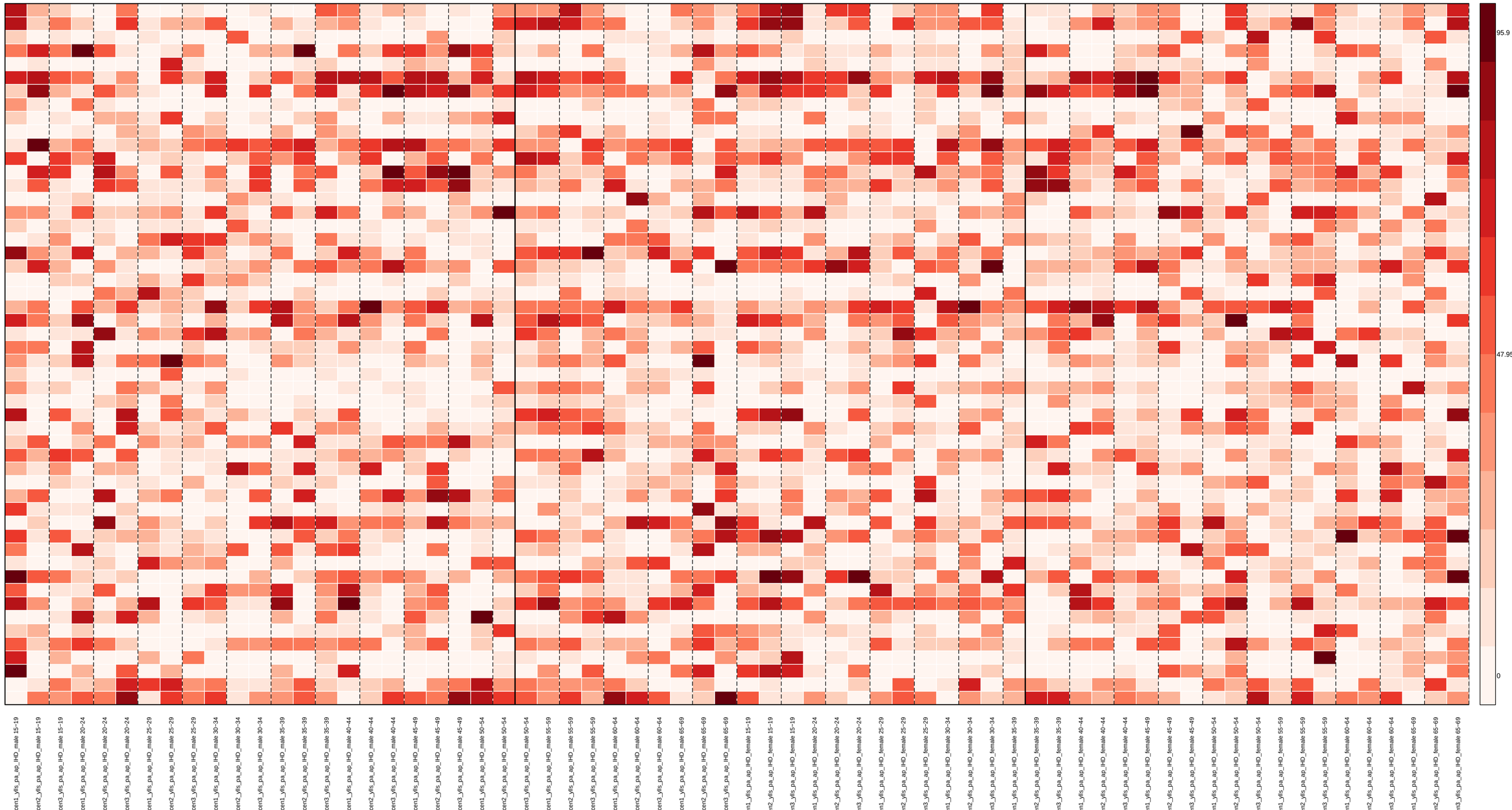
copiapo − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



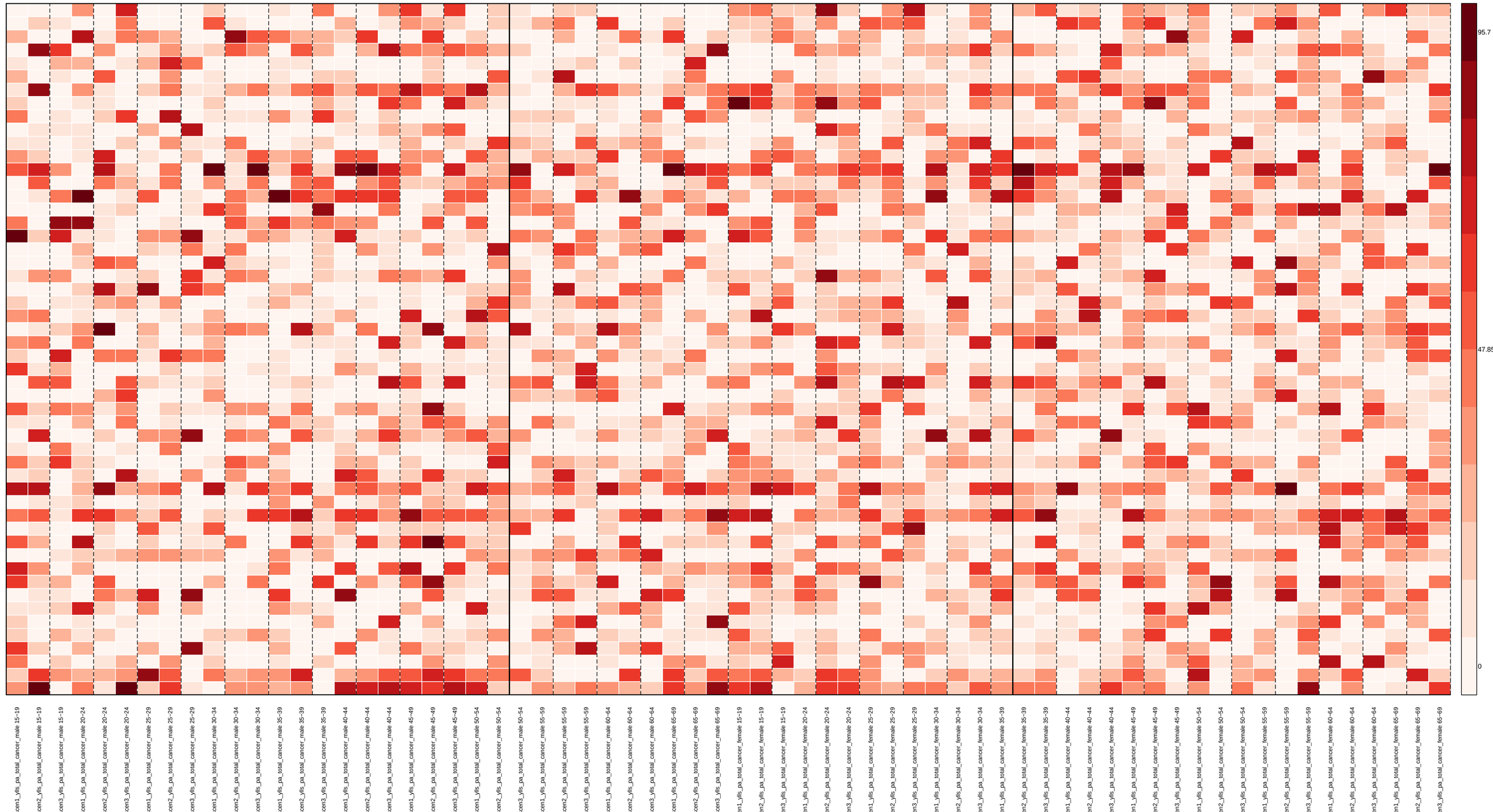
coquimbo_laserena − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



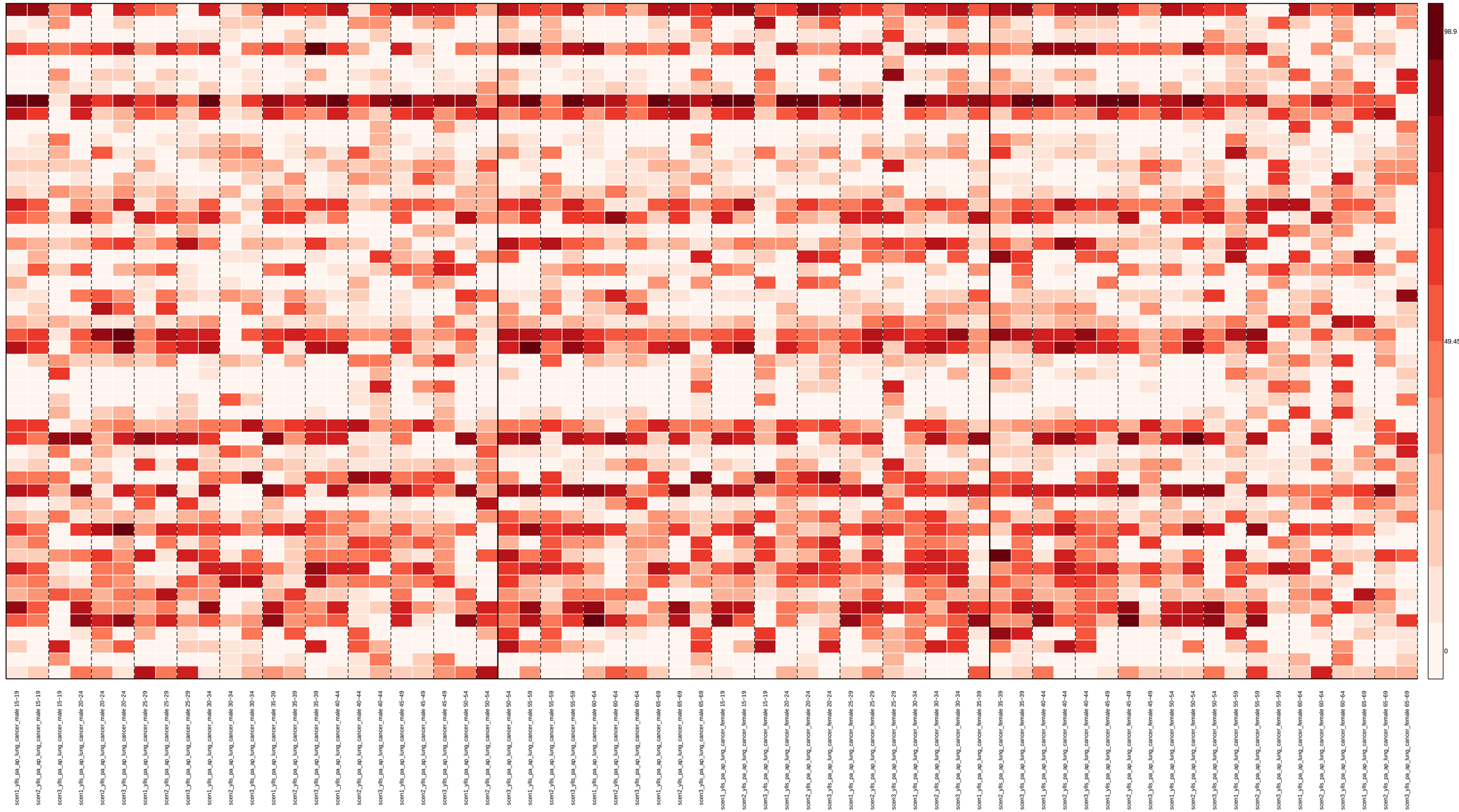
coquimbo_laserena − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



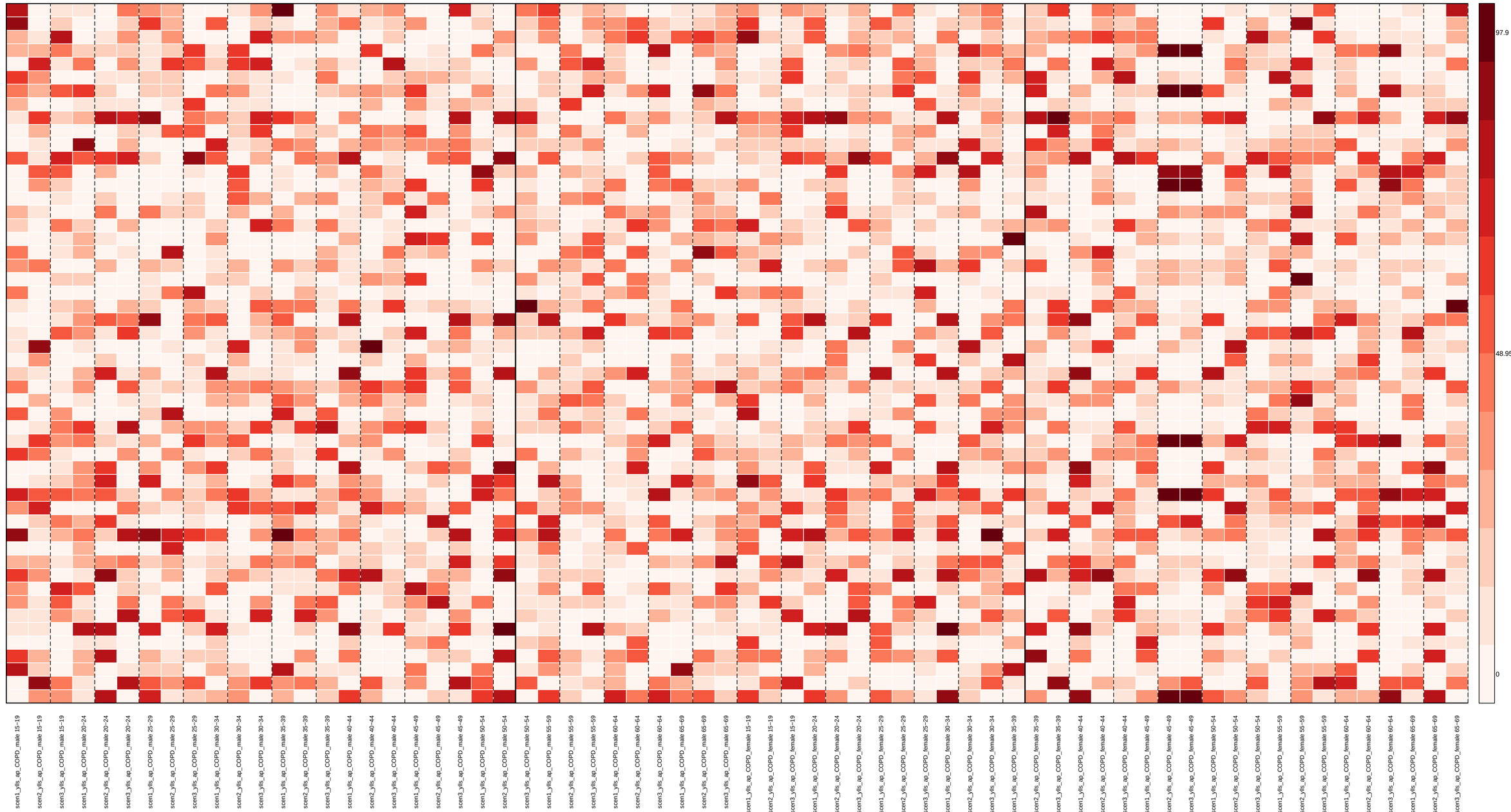
coquimbo_laserena − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



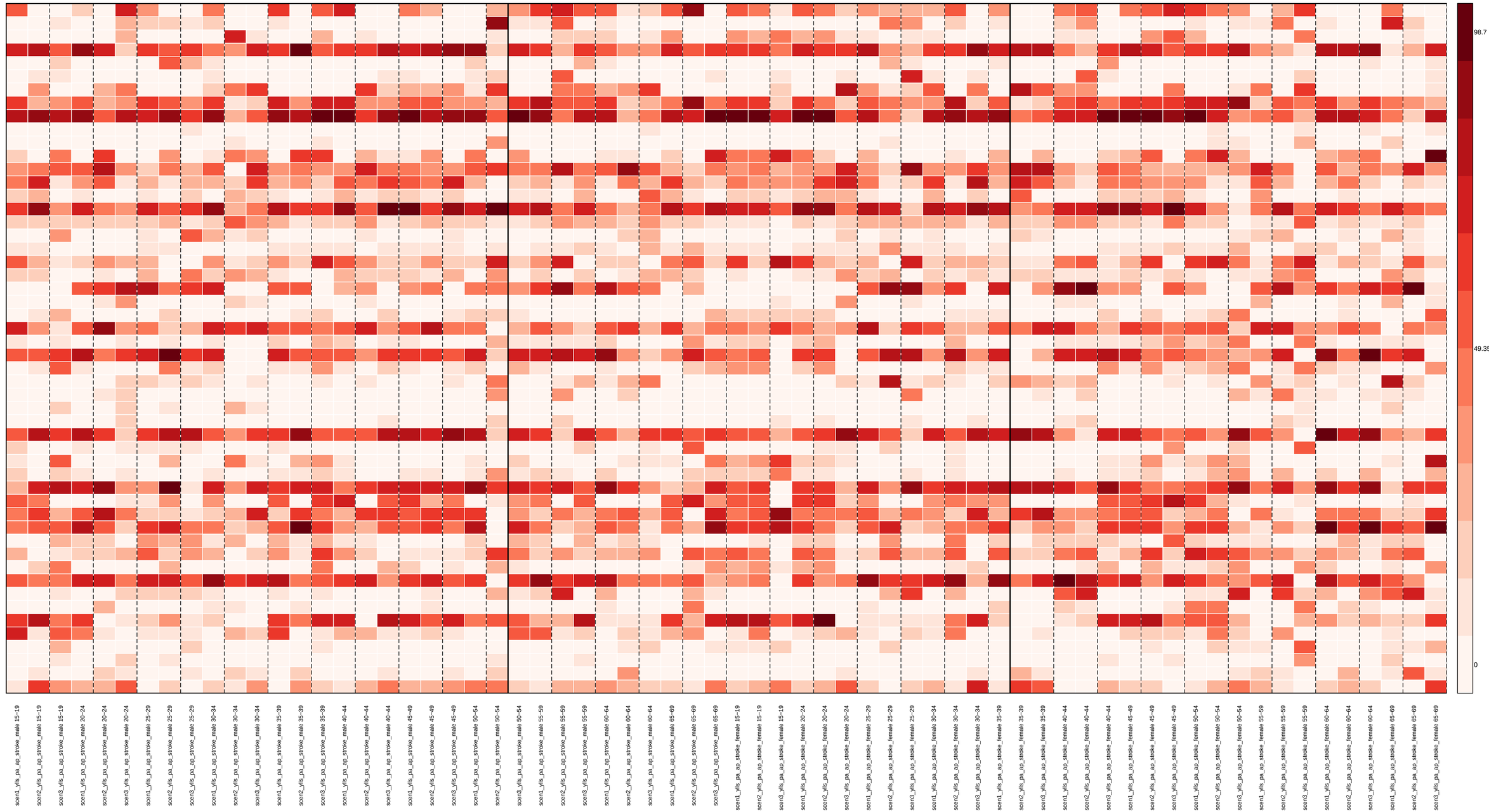
coquimbo_laserena − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



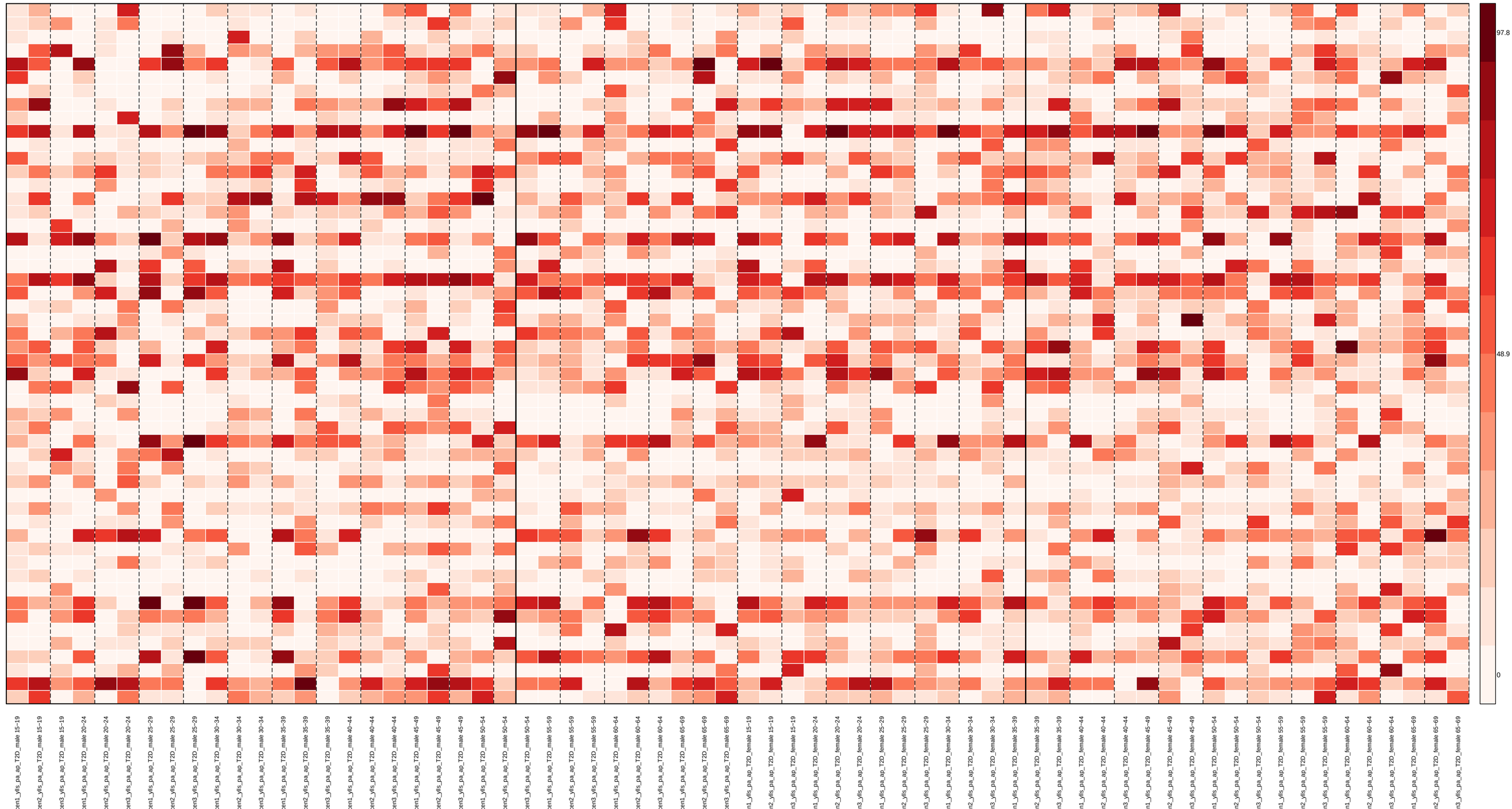
coquimbo_laserena − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



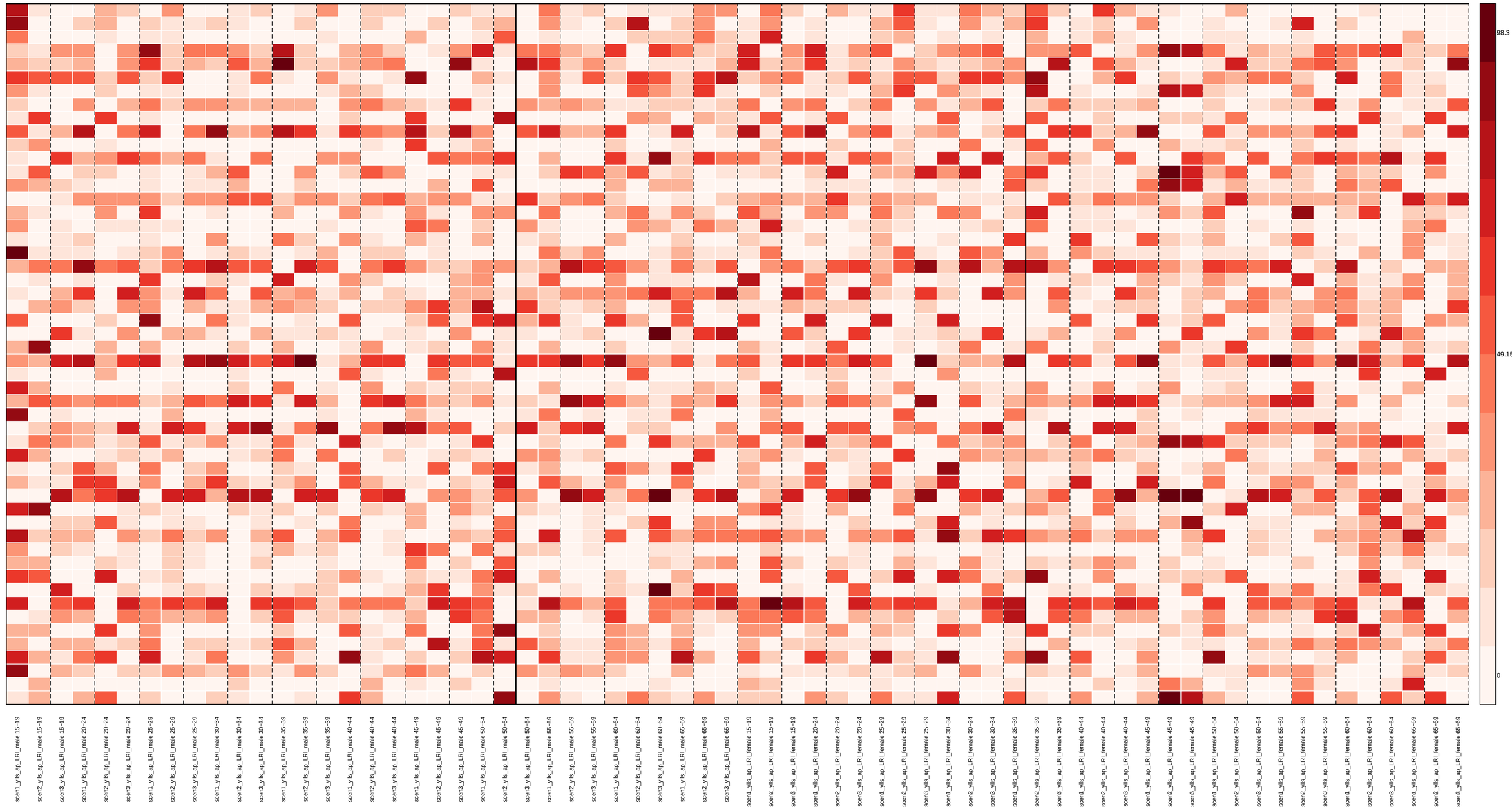
coquimbo_laserena − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



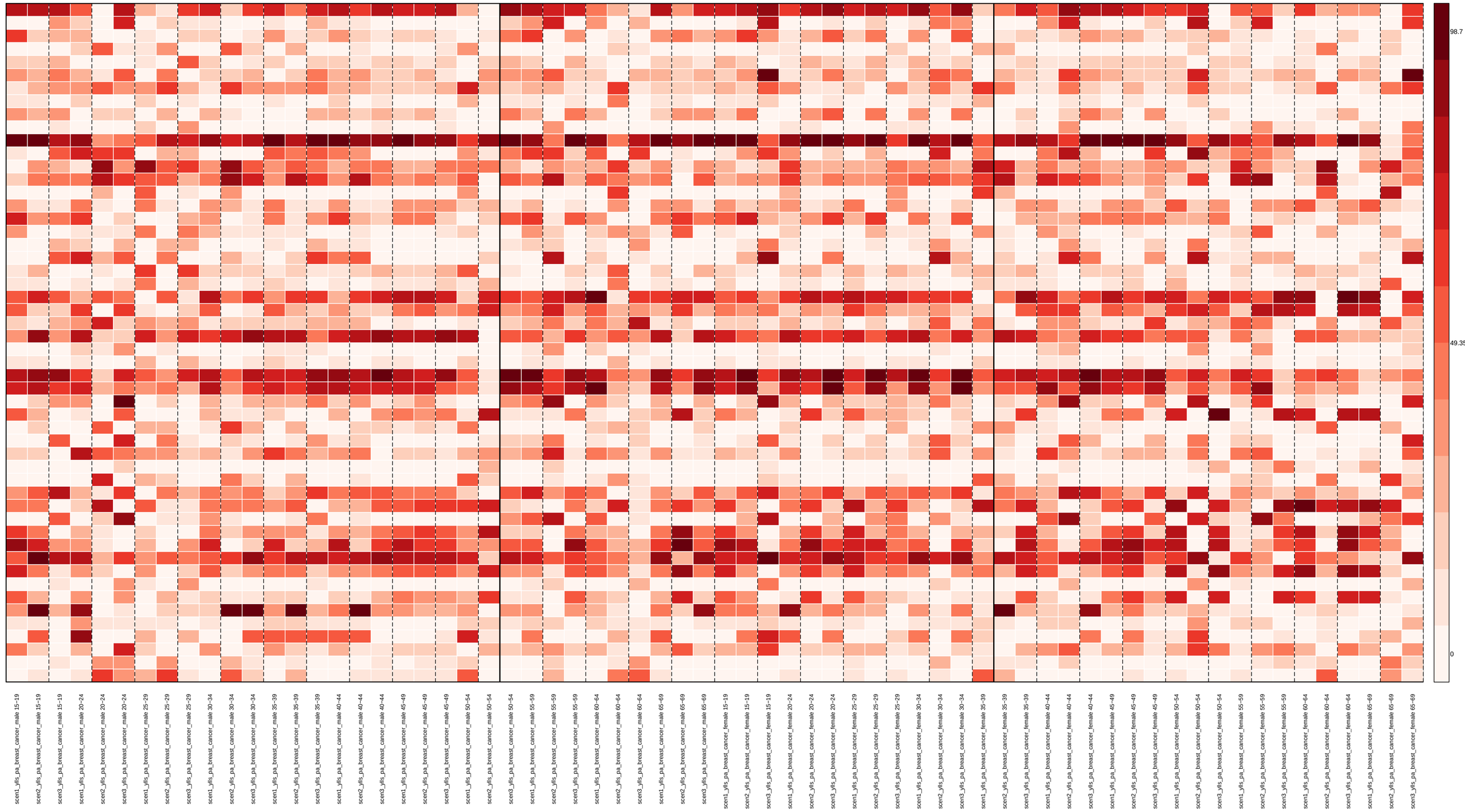
coquimbo_laserena − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



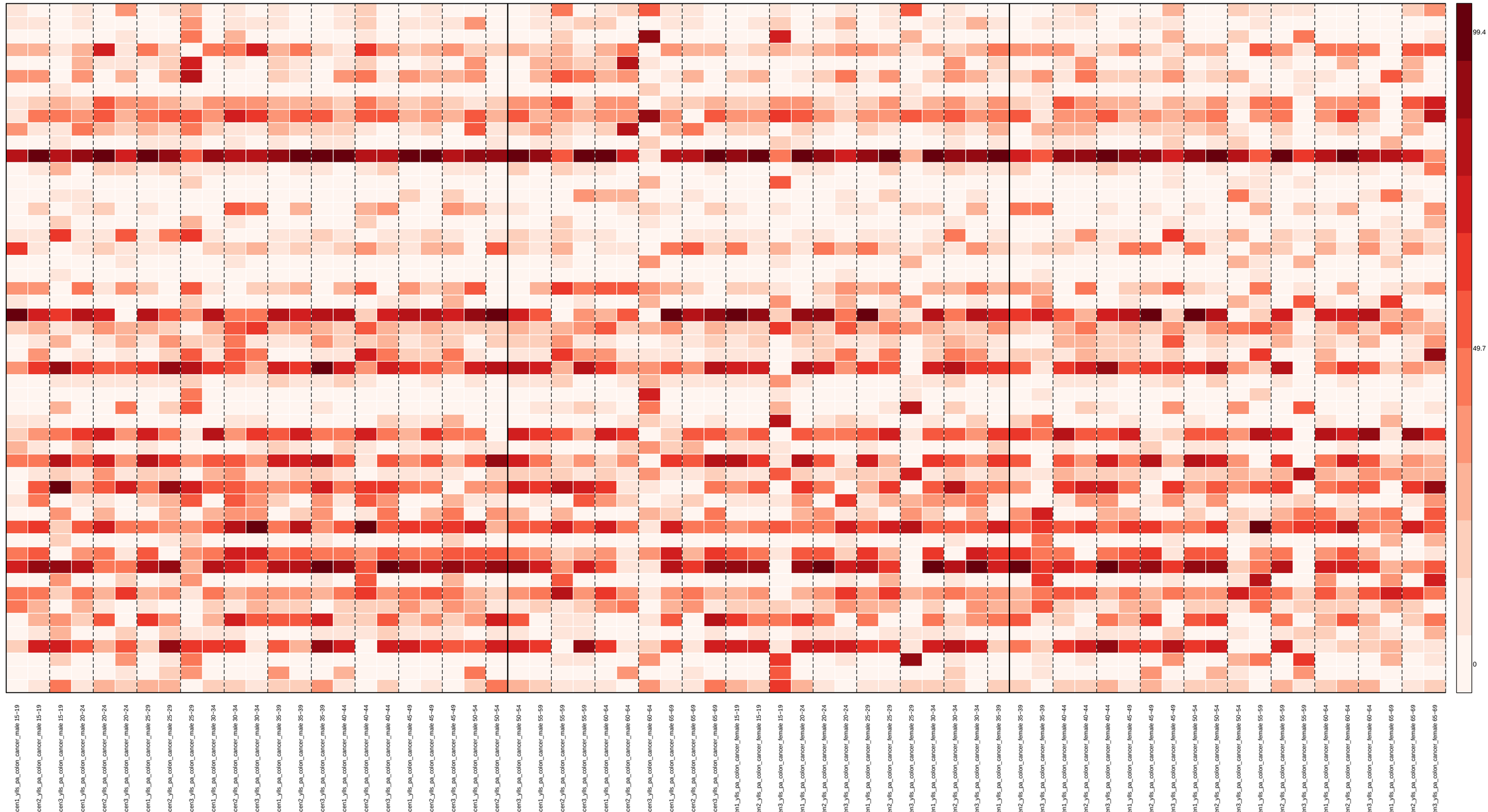
coquimbo_laserena − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



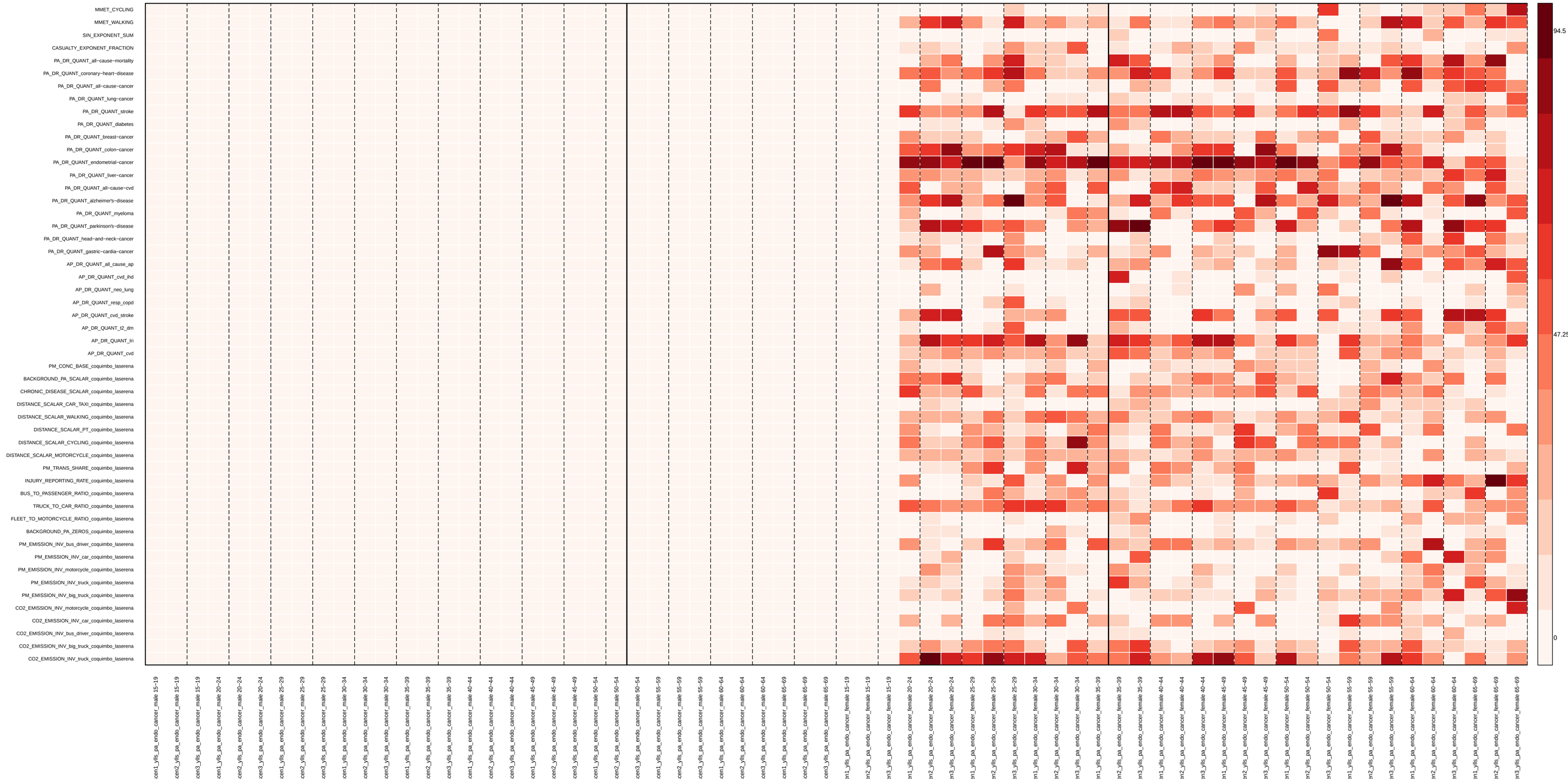
coquimbo_laserena − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



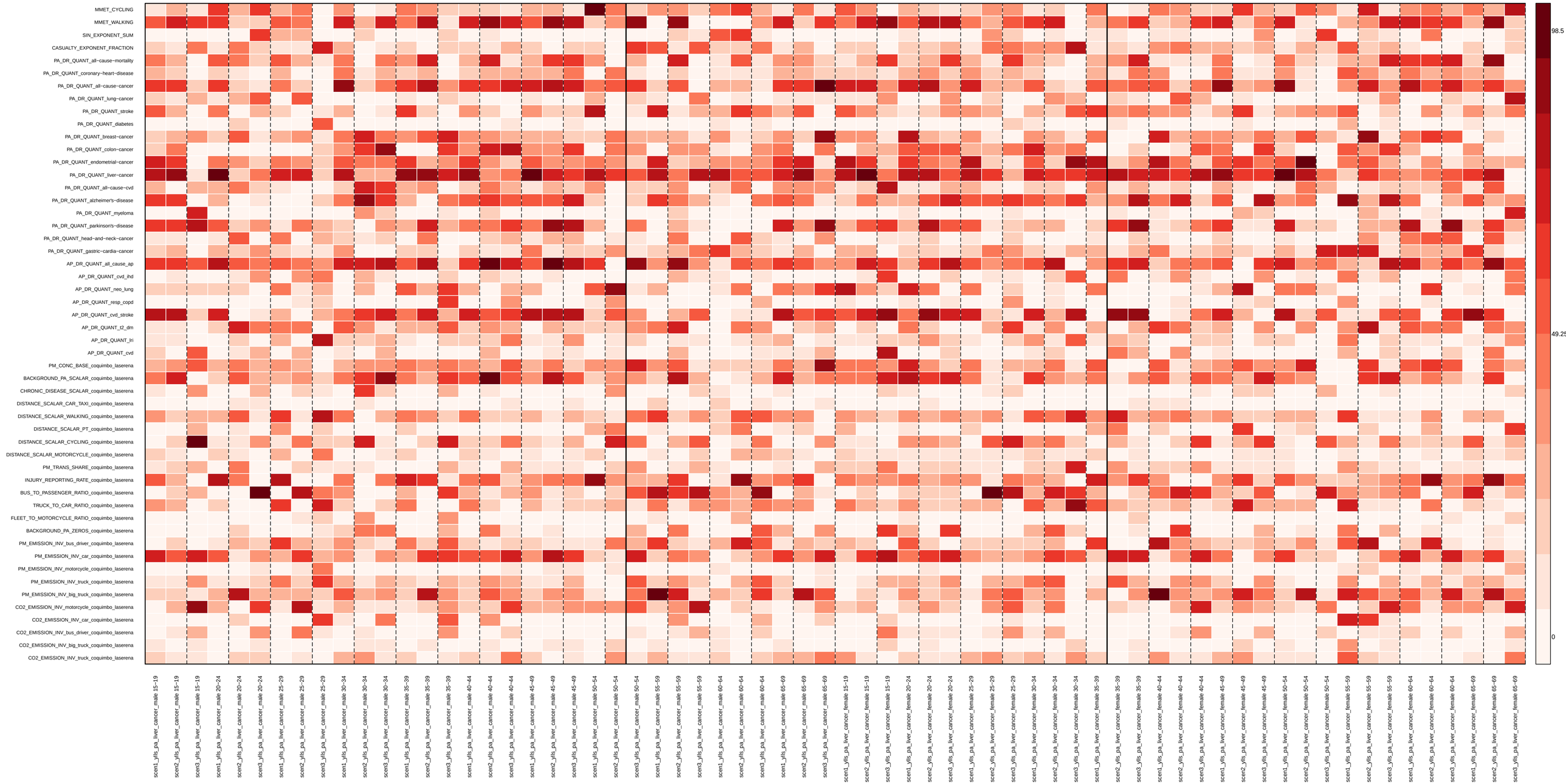
coquimbo_laserena − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



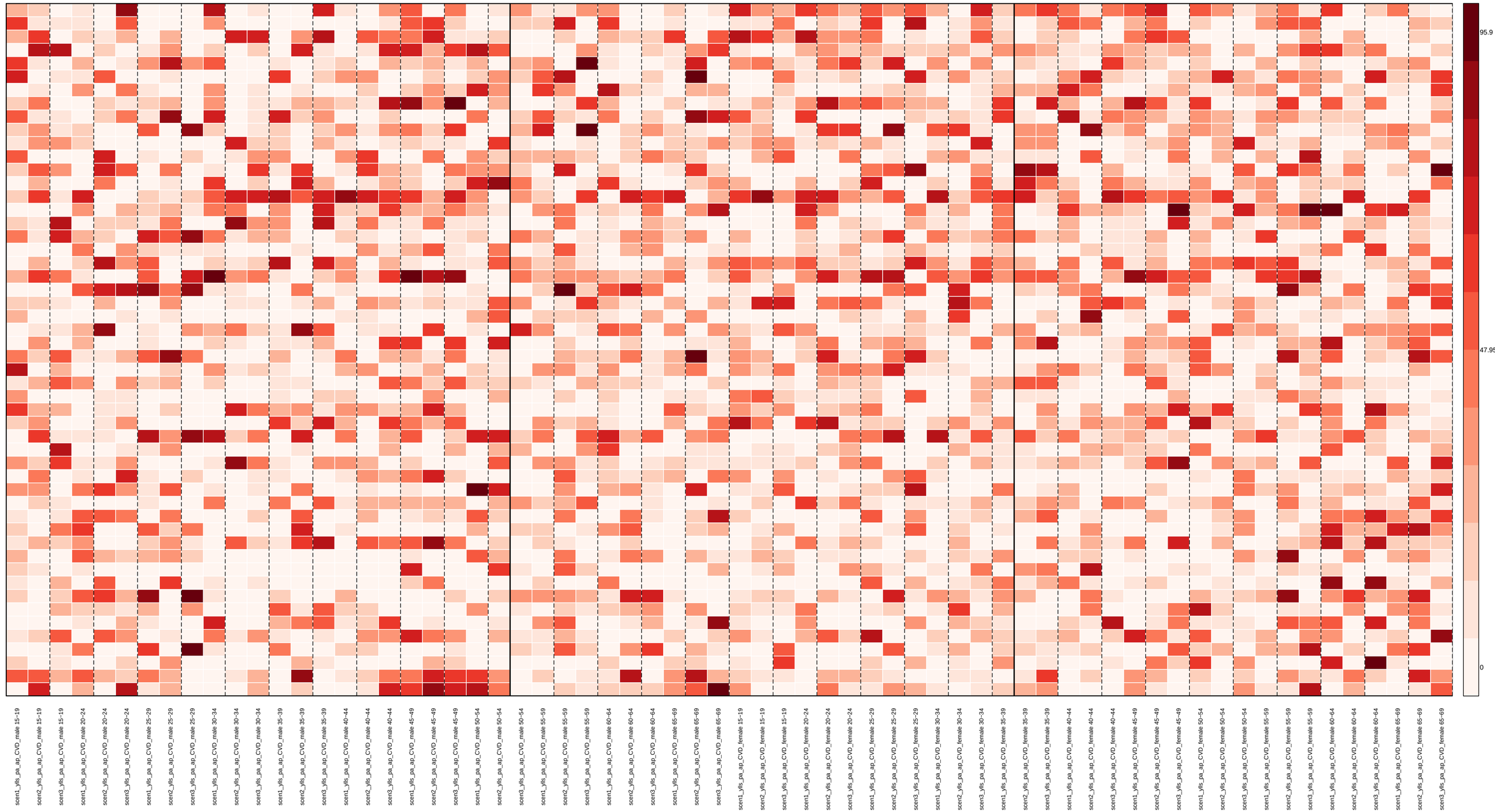
coquimbo_laserena − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



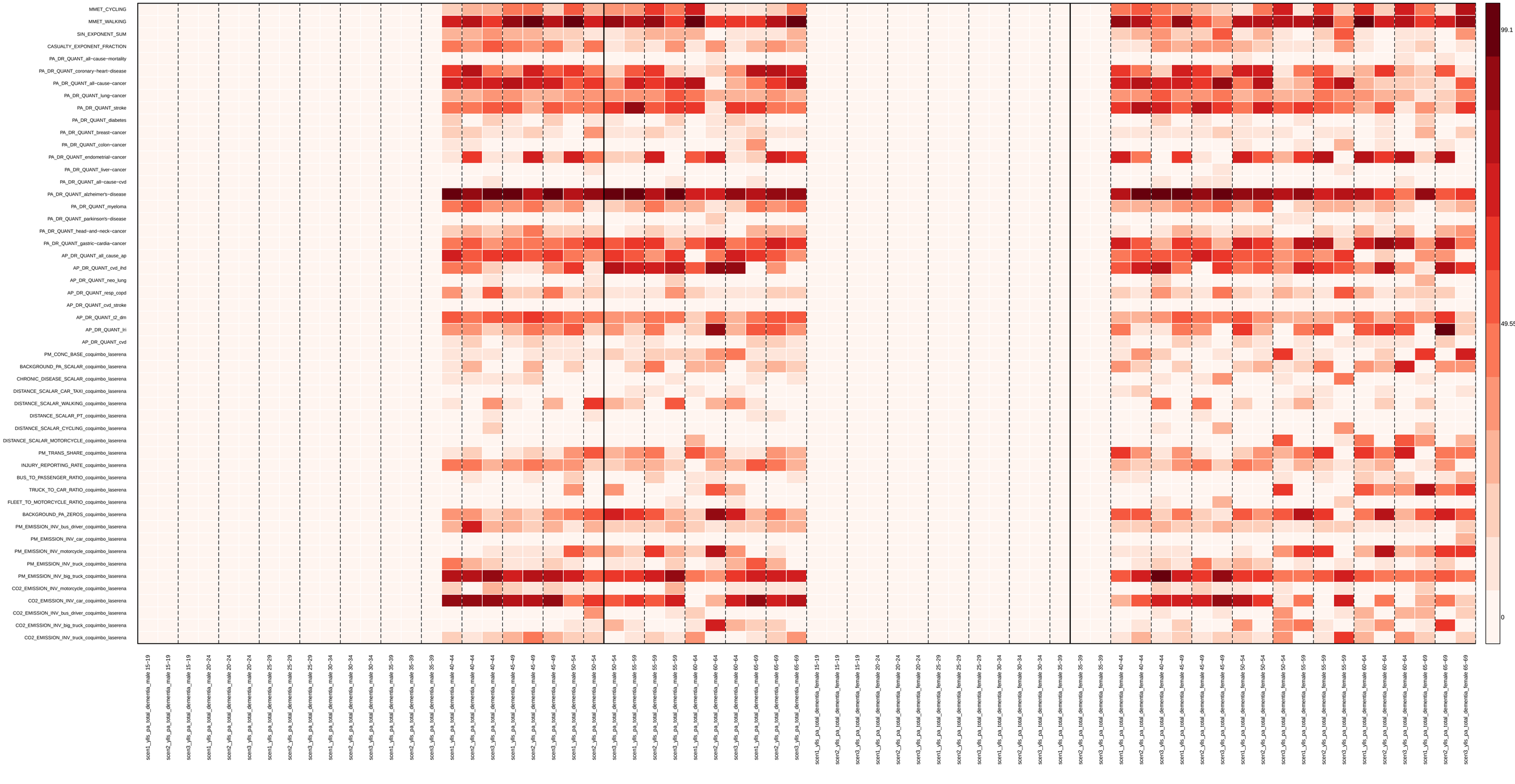
coquimbo_laserena − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



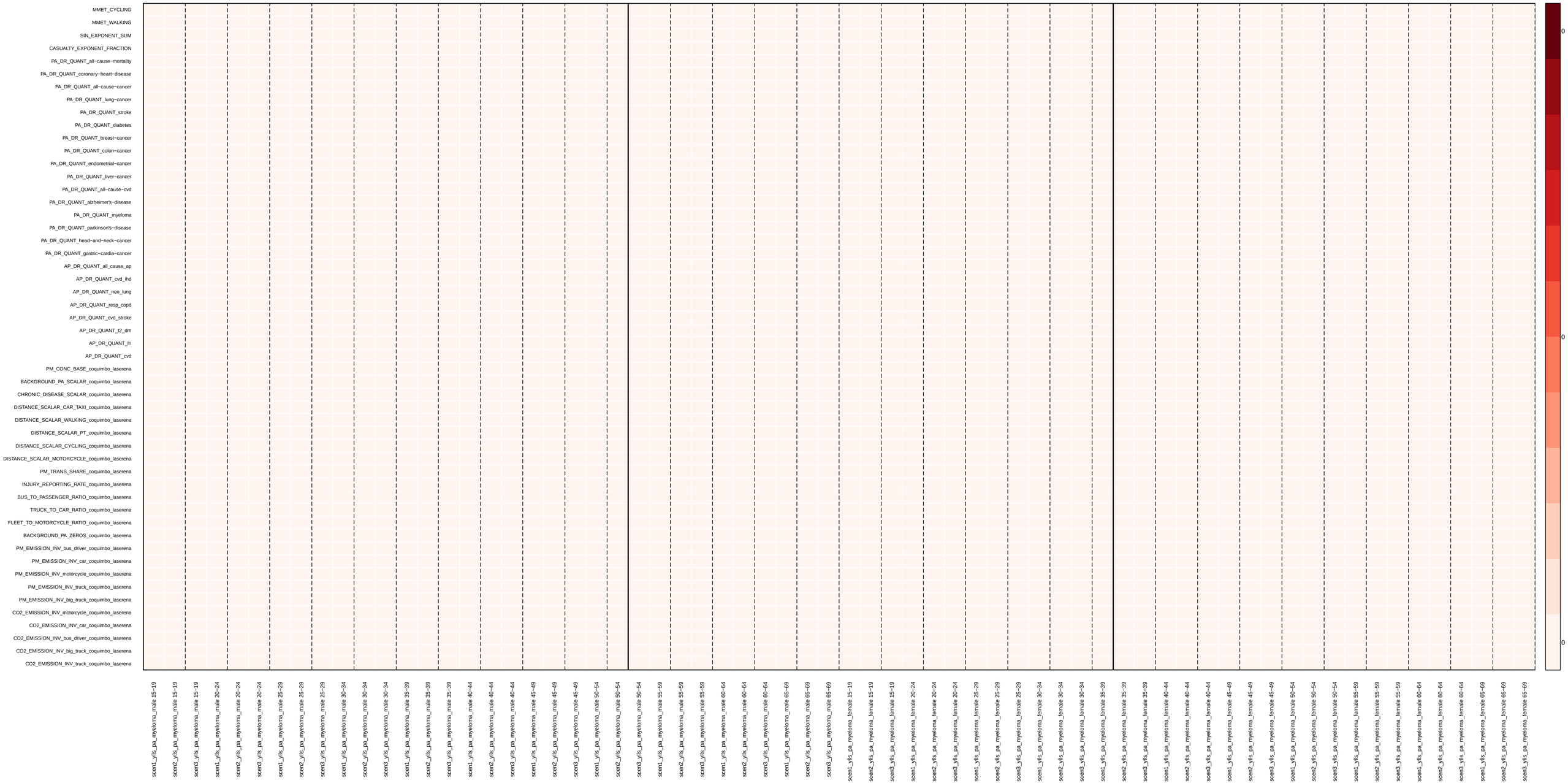
coquimbo_laserena − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



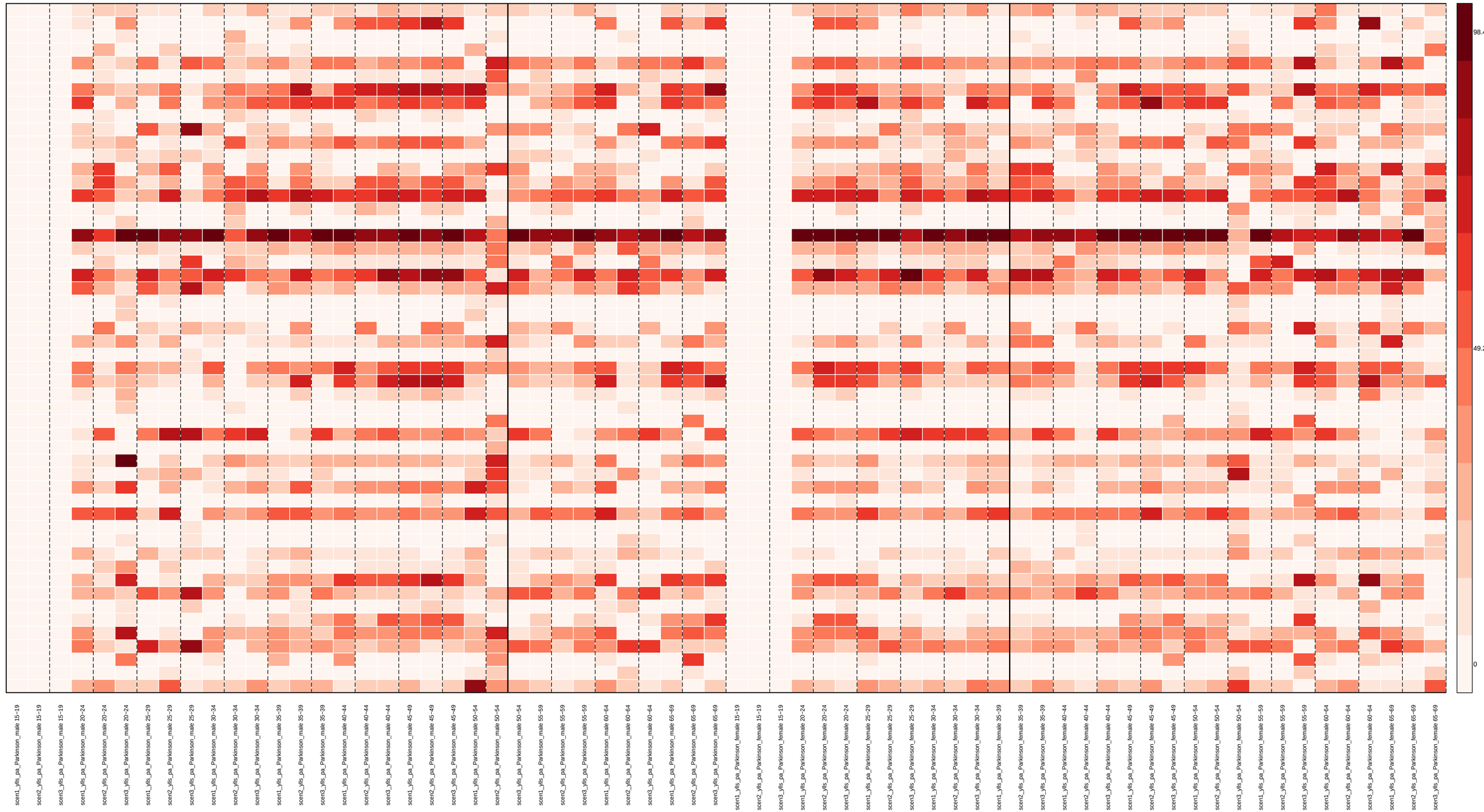
coquimbo_laserena − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



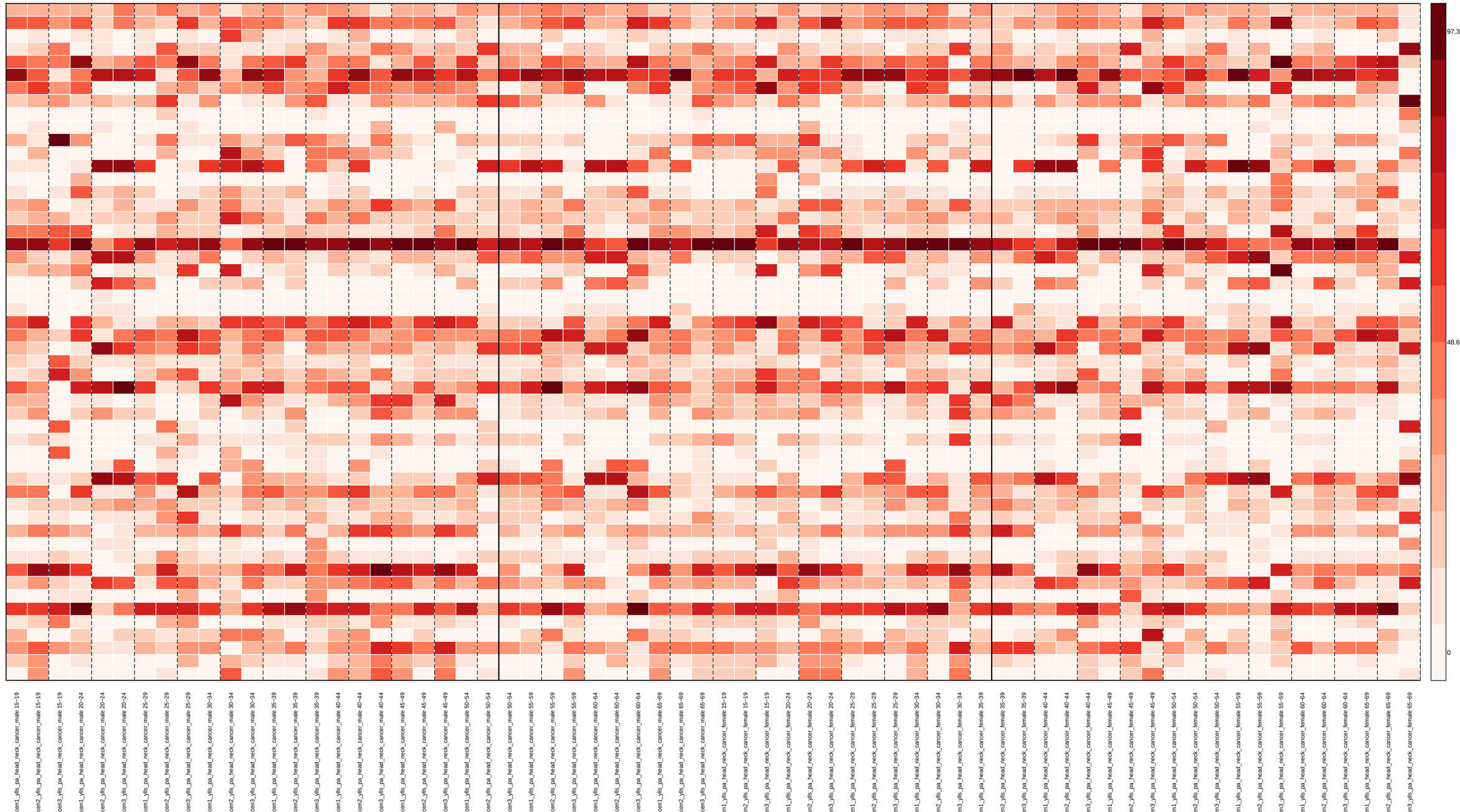
coquimbo_laserena − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



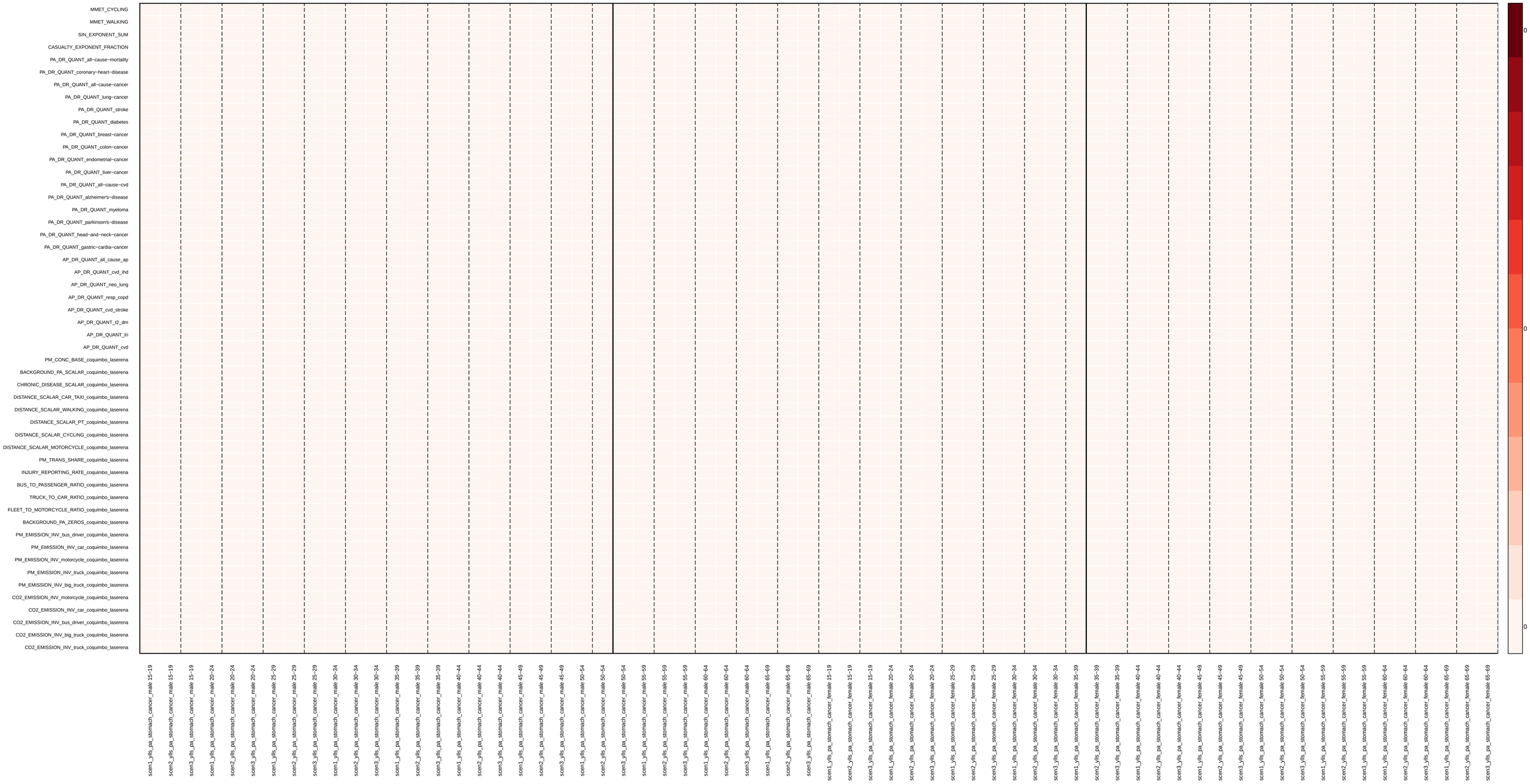
coquimbo_laserena − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



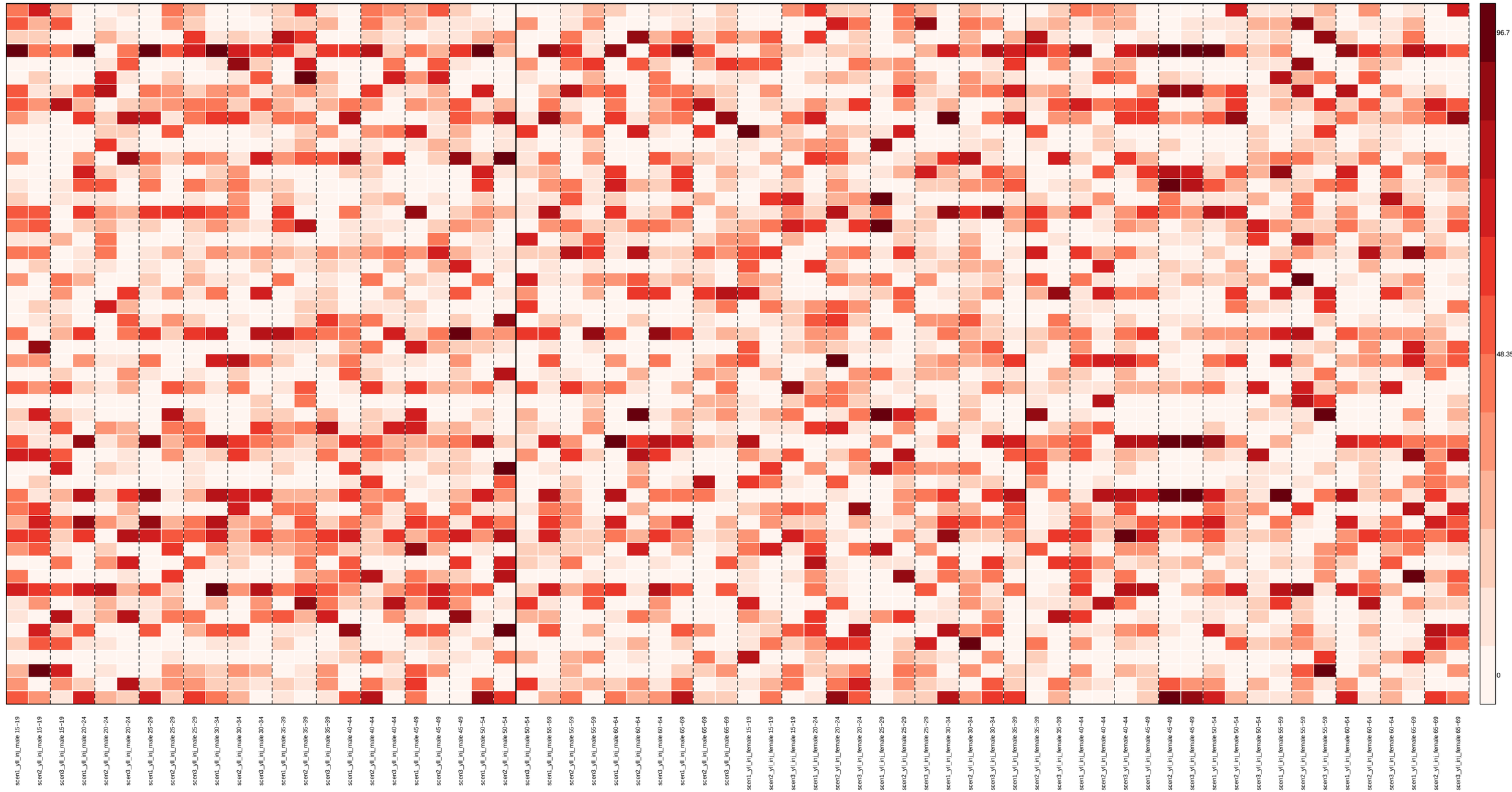
coquimbo_laserena − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



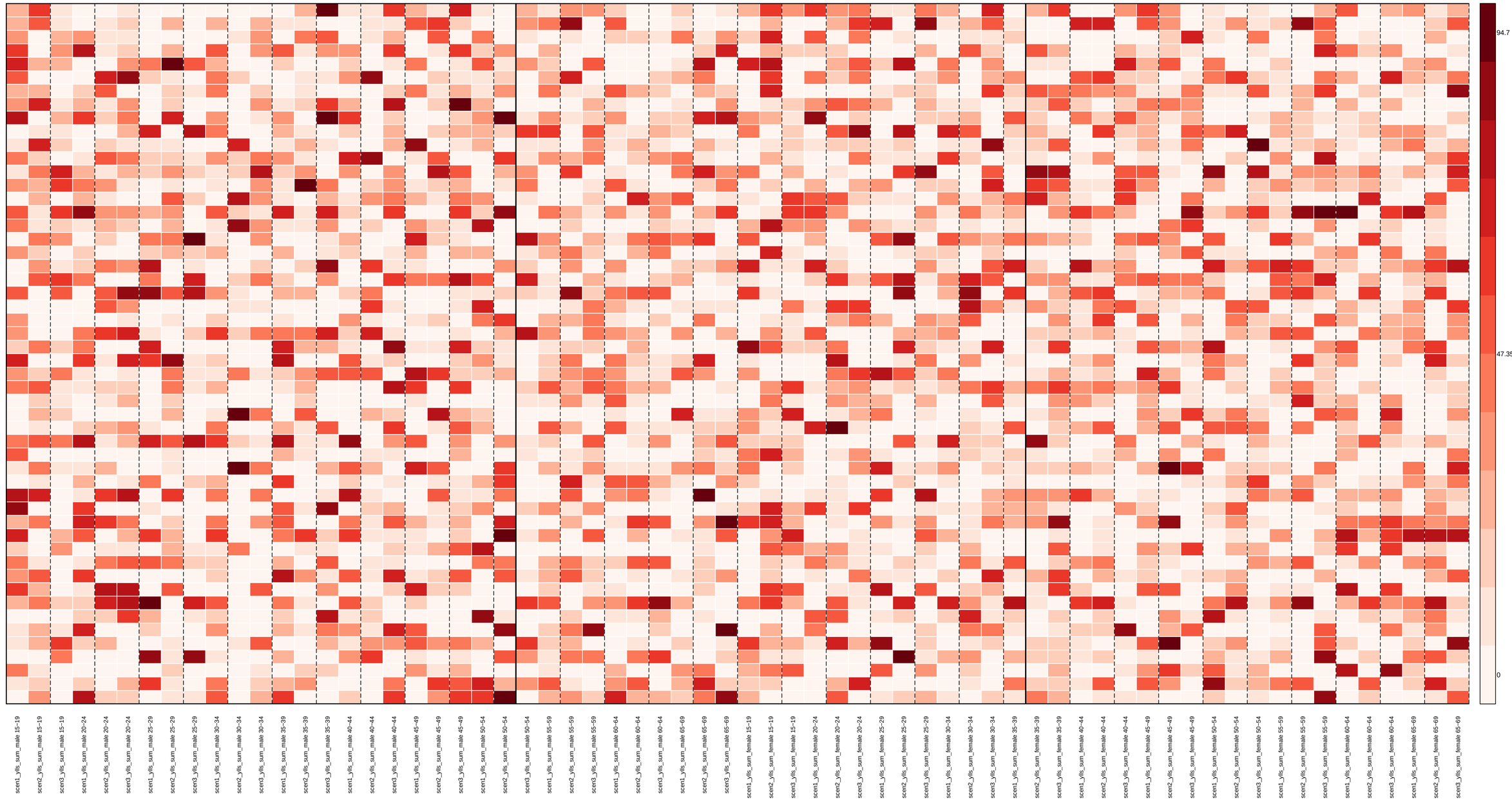
coquimbo_laserena − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



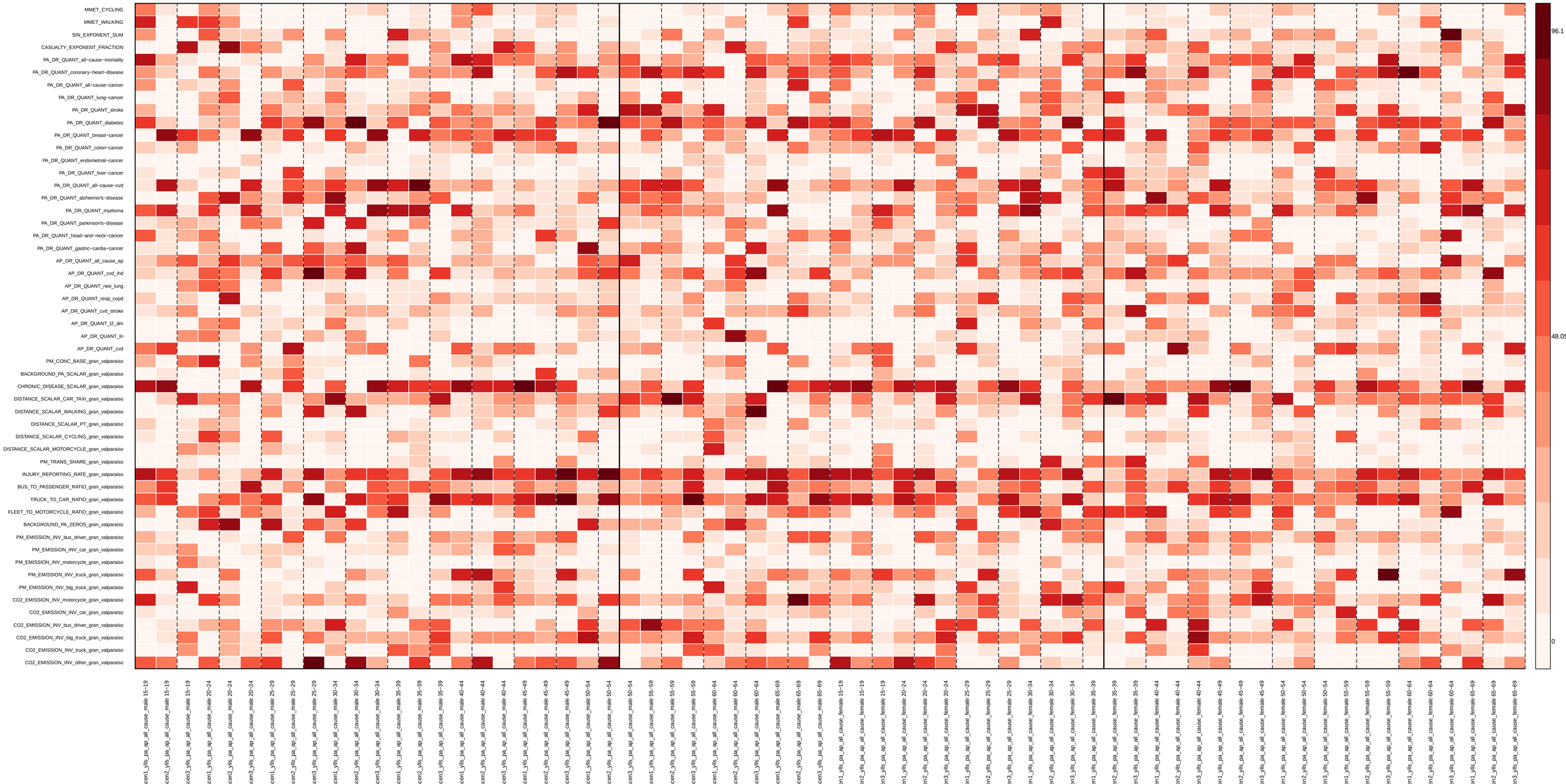
coquimbo_laserena − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



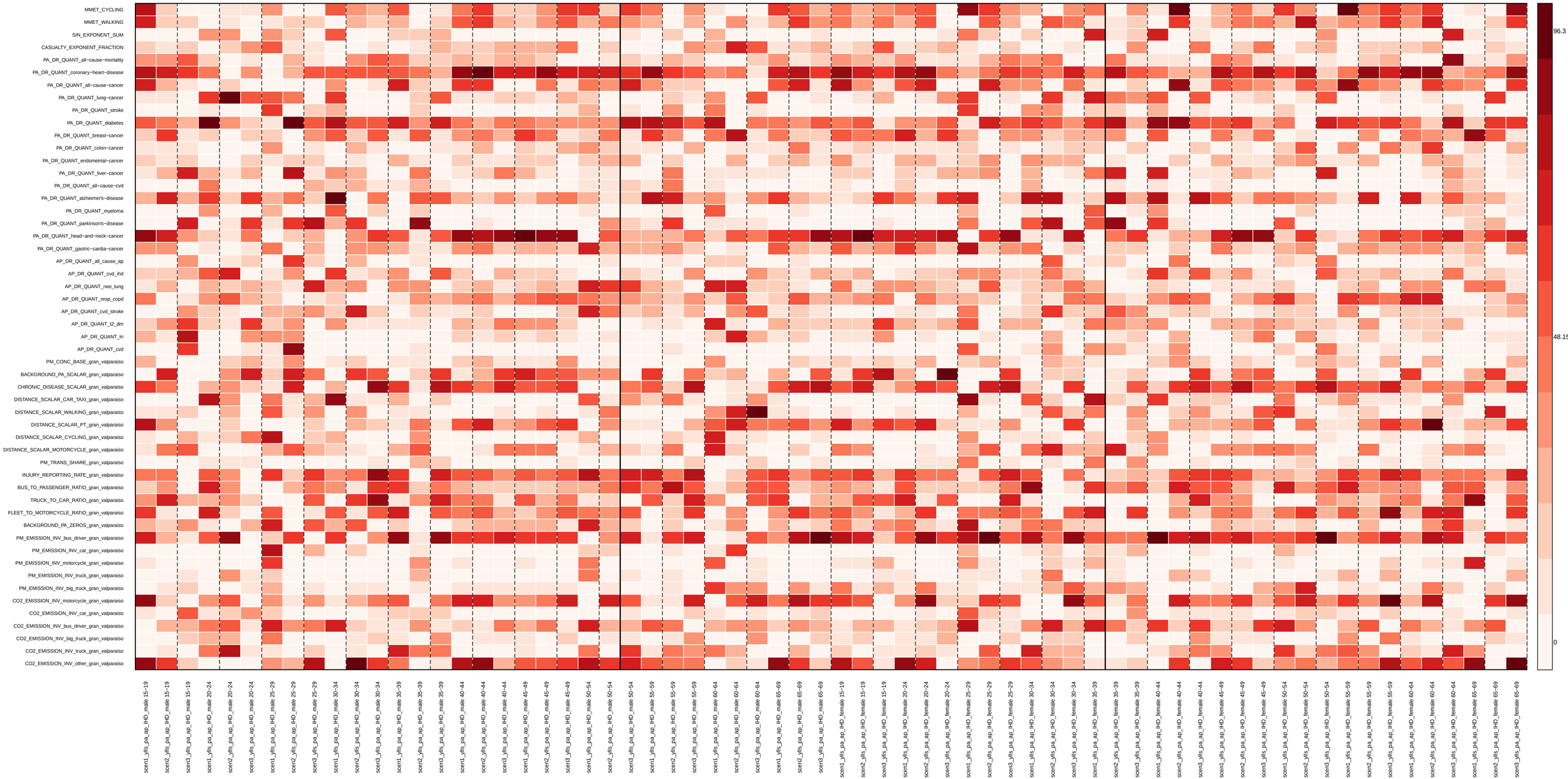
coquimbo_laserena − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



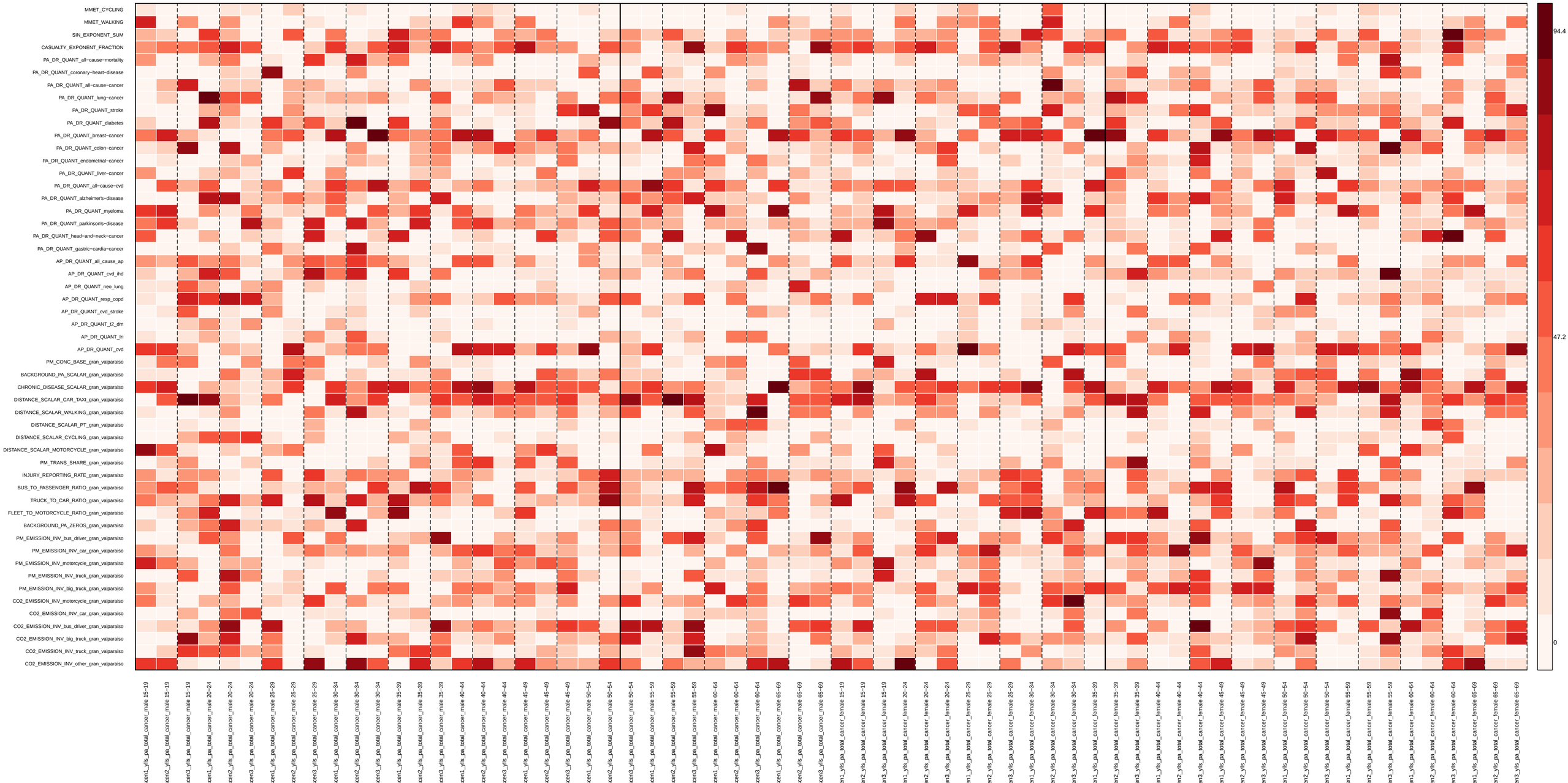
gran_valparaiso − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



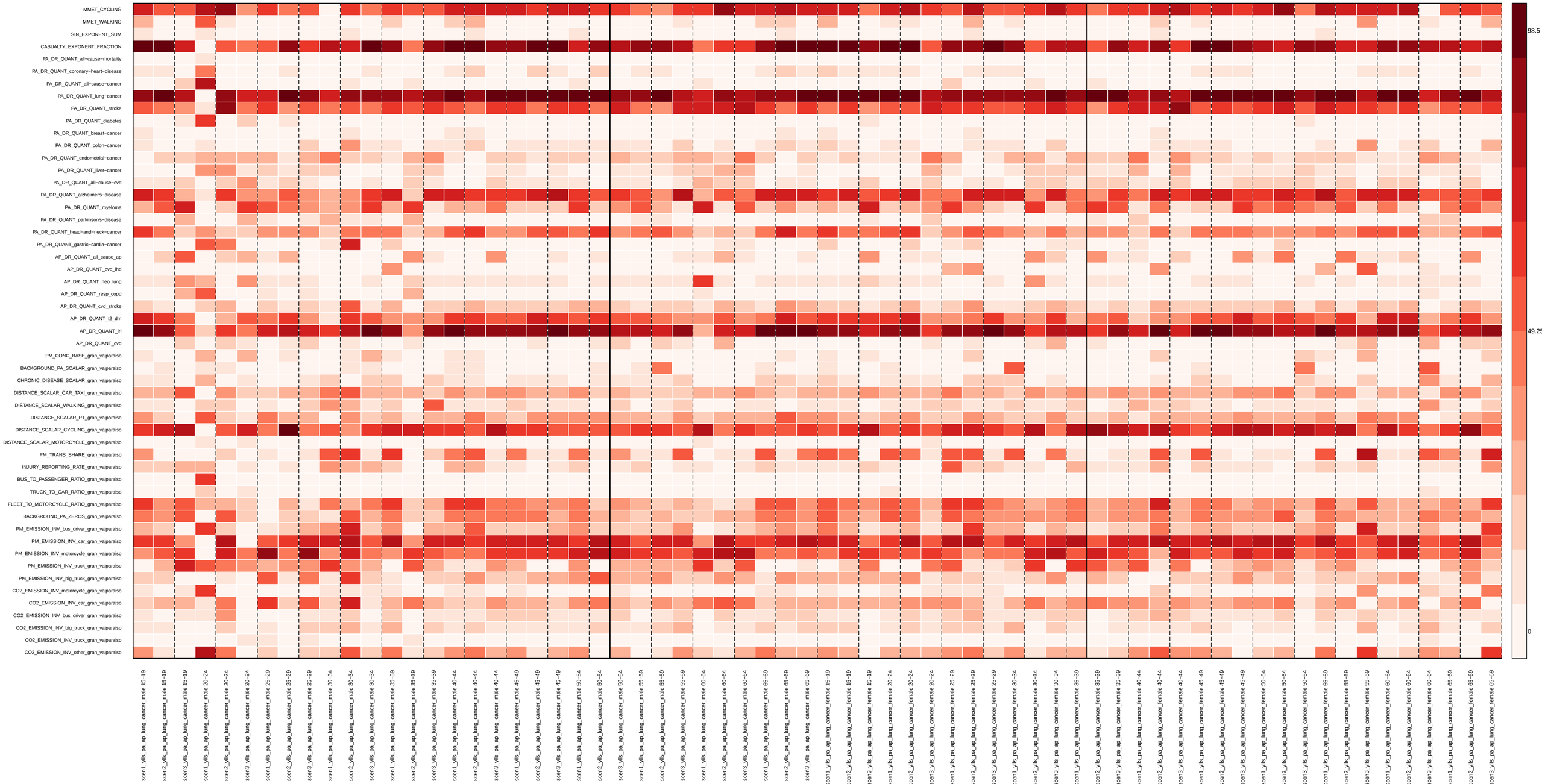
gran_valparaiso − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



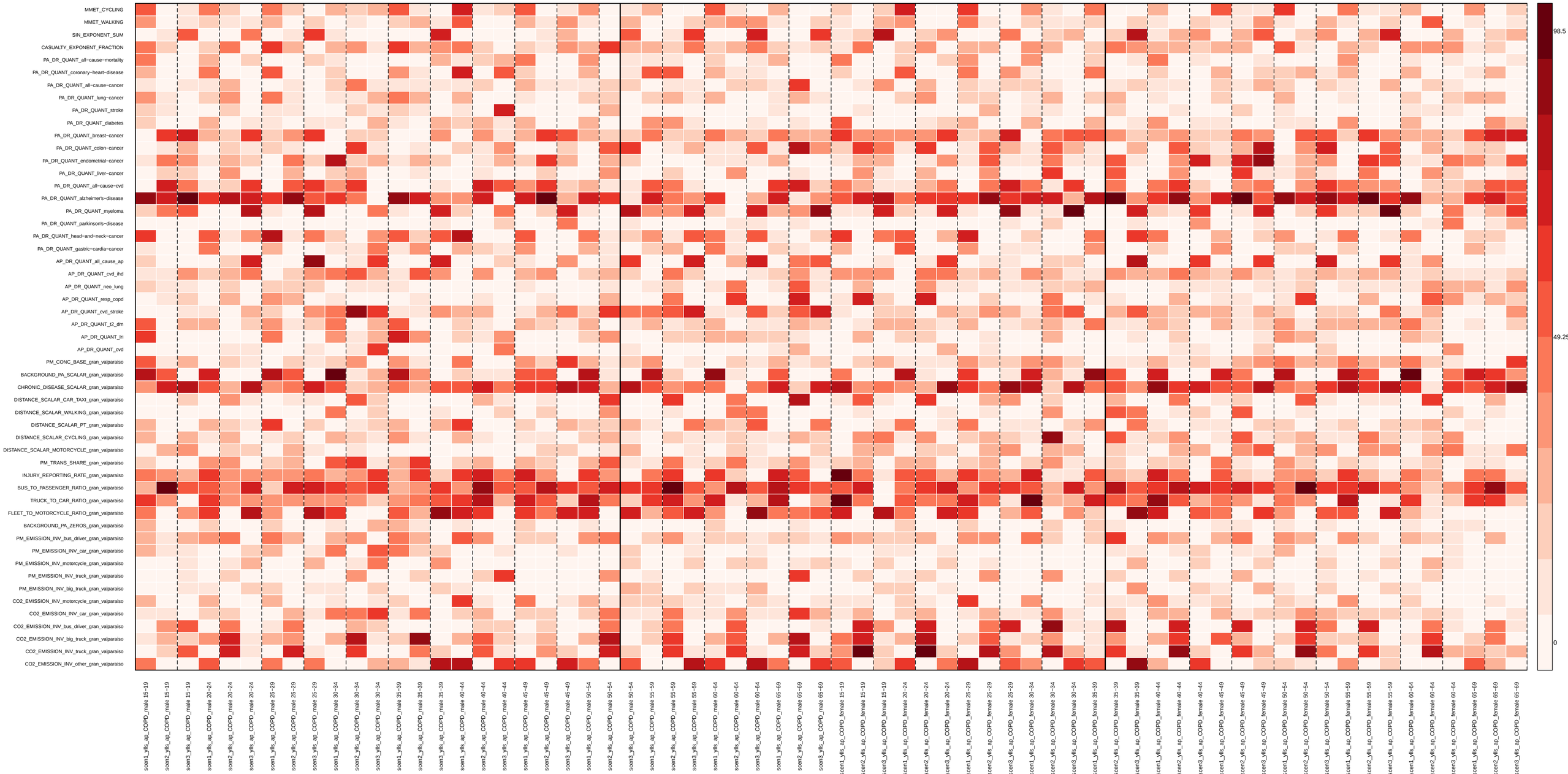
gran_valparaiso − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



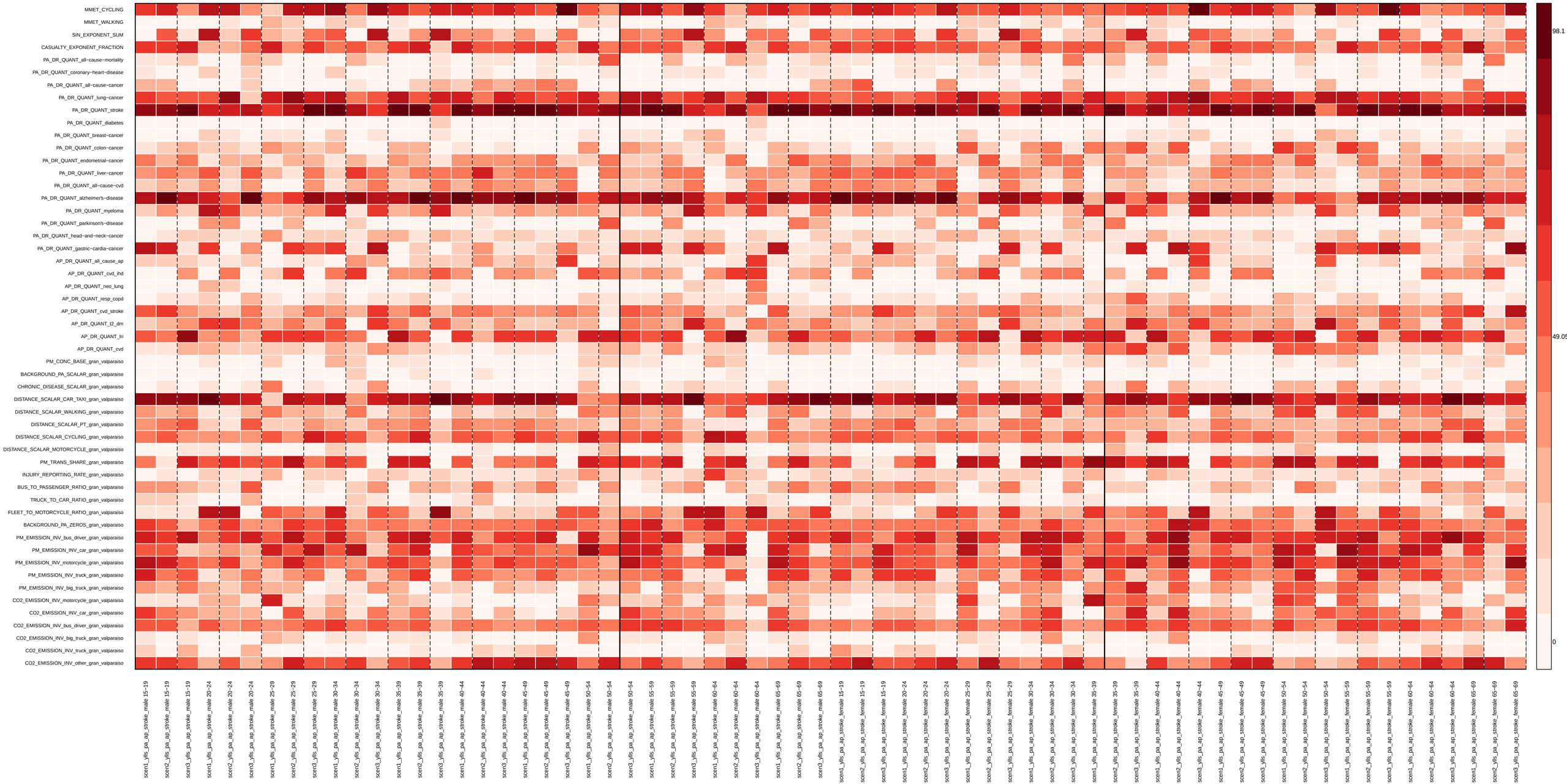
gran_valparaiso − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



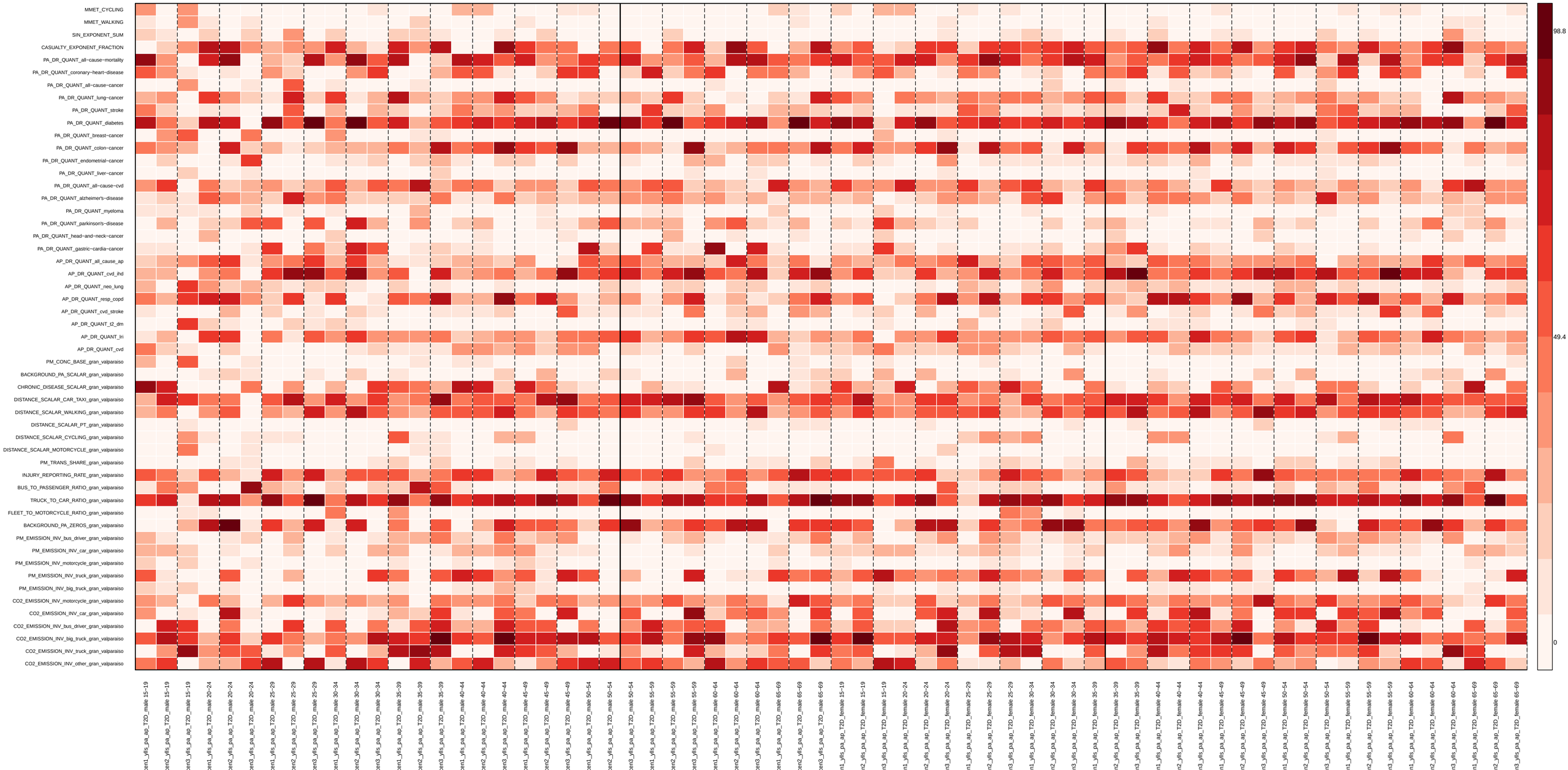
gran_valparaiso − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



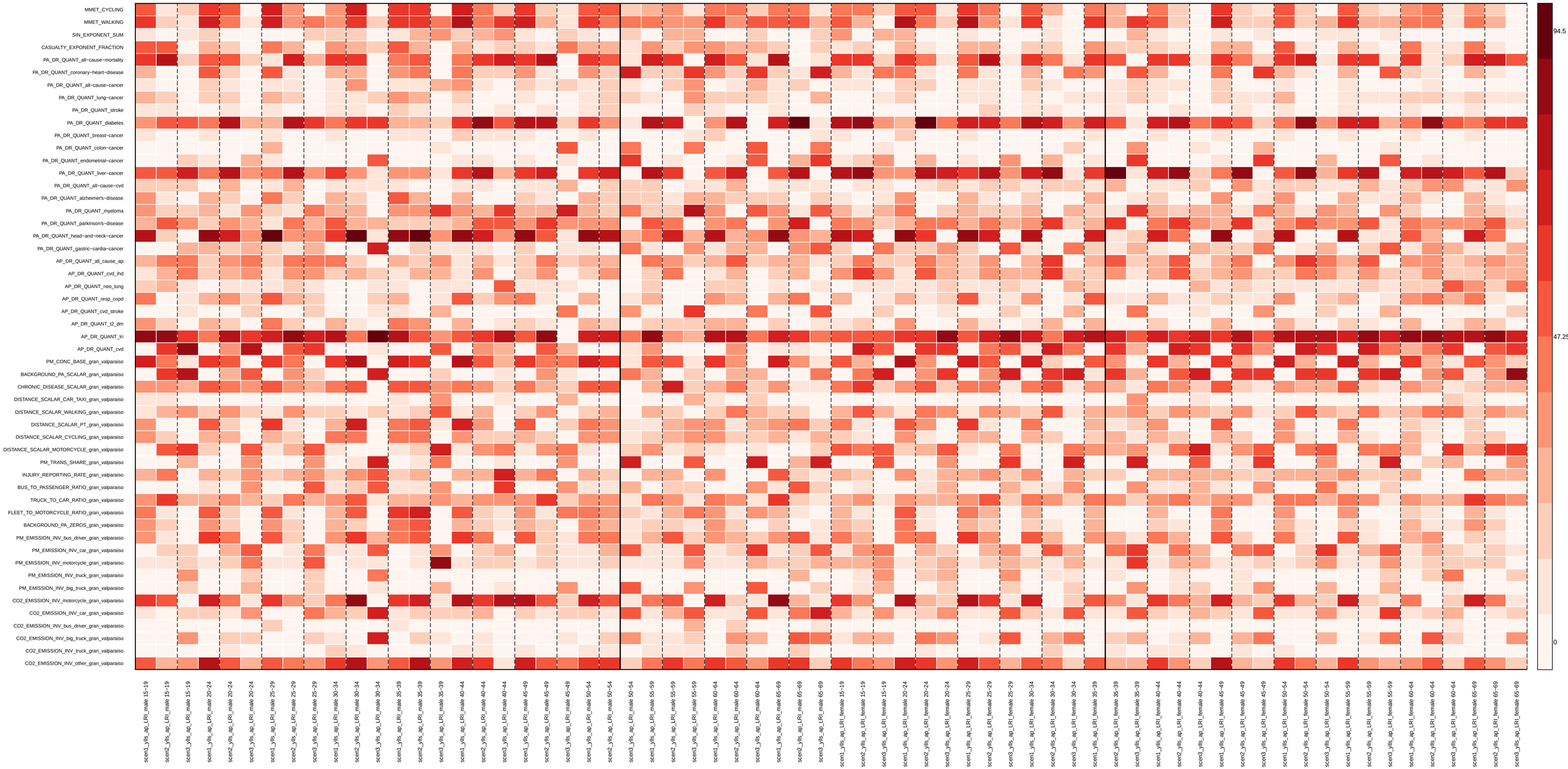
gran_valparaiso − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



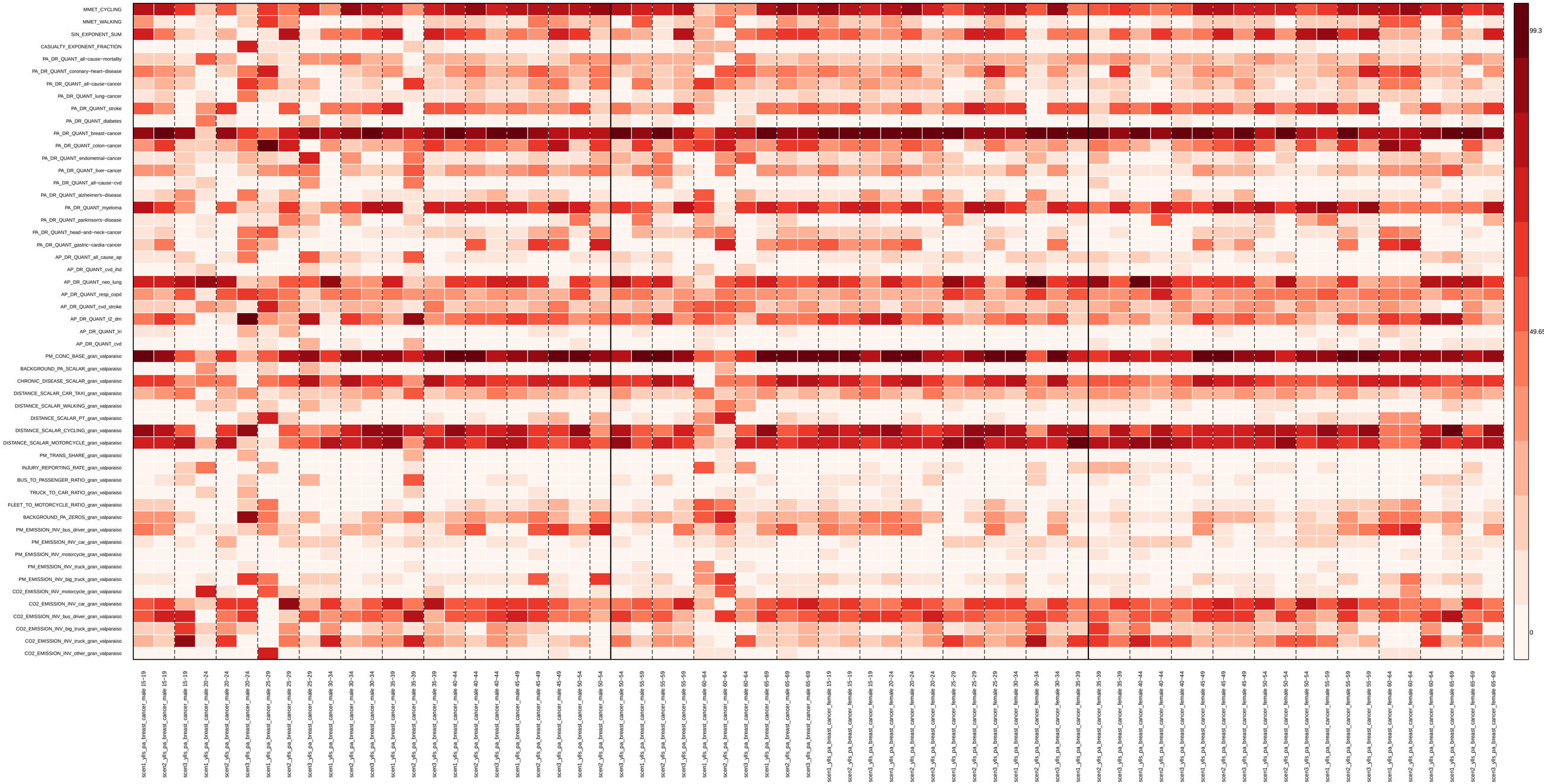
gran_valparaiso − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



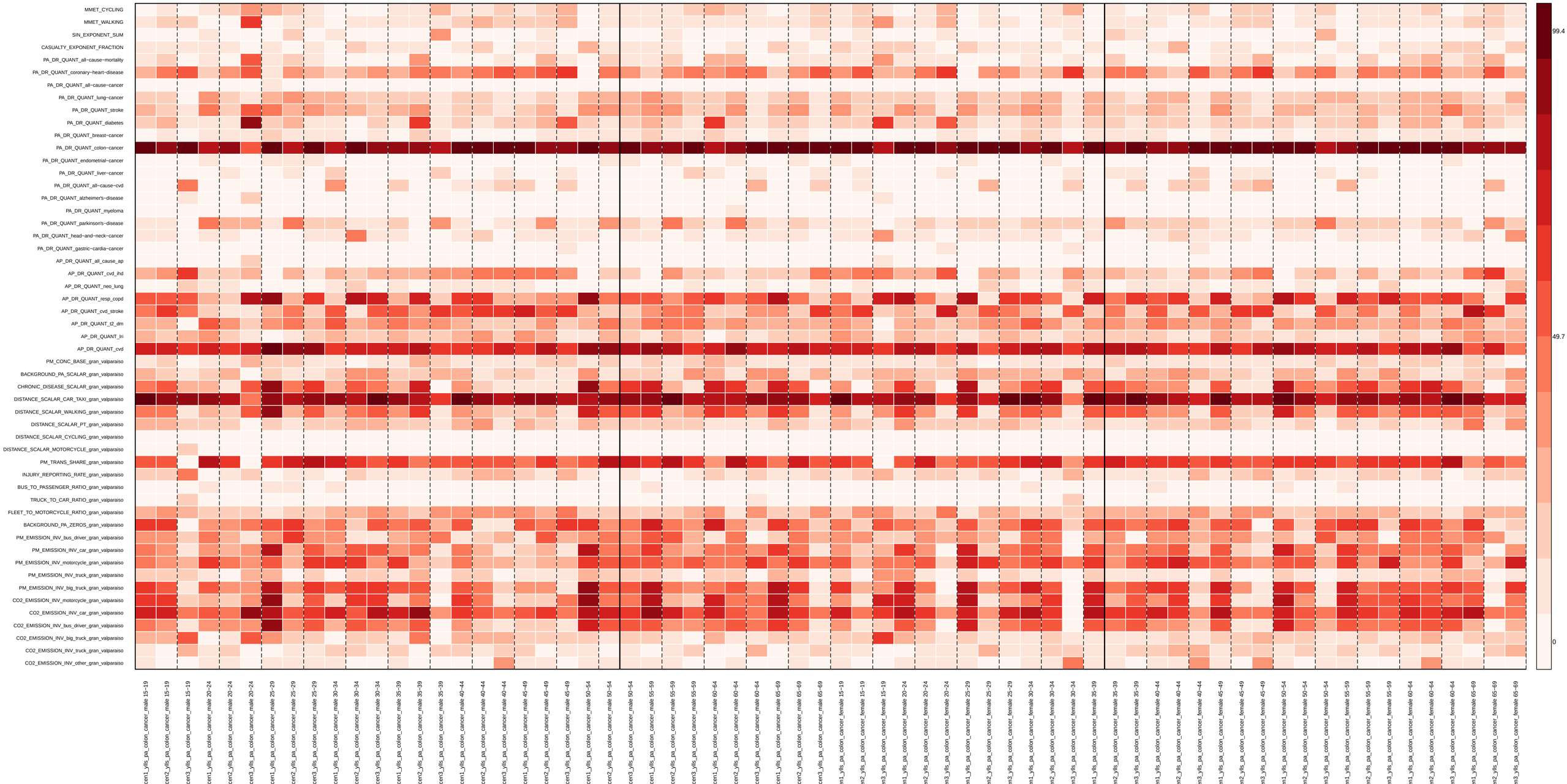
gran_valparaiso − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



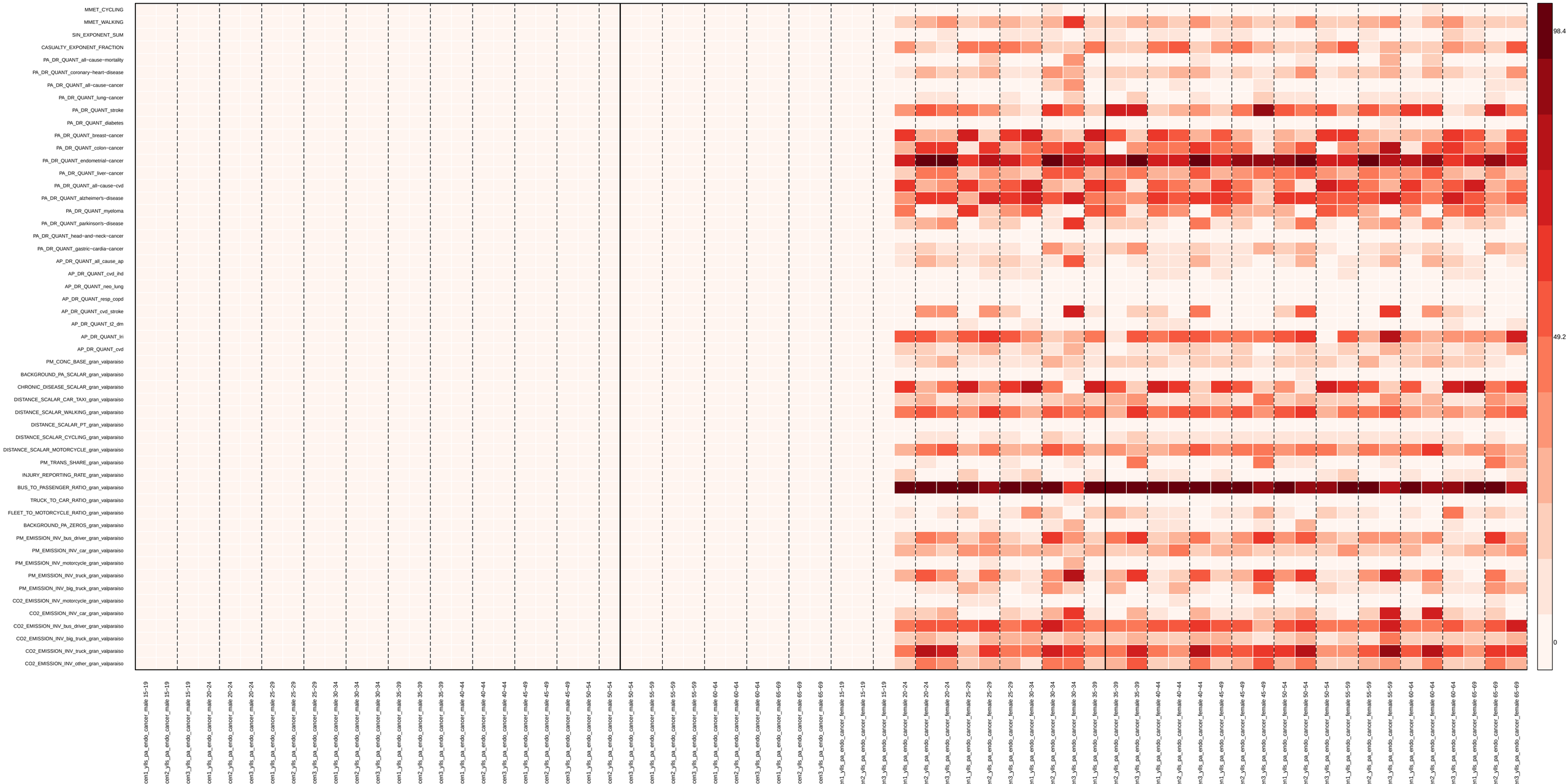
gran_valparaiso − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



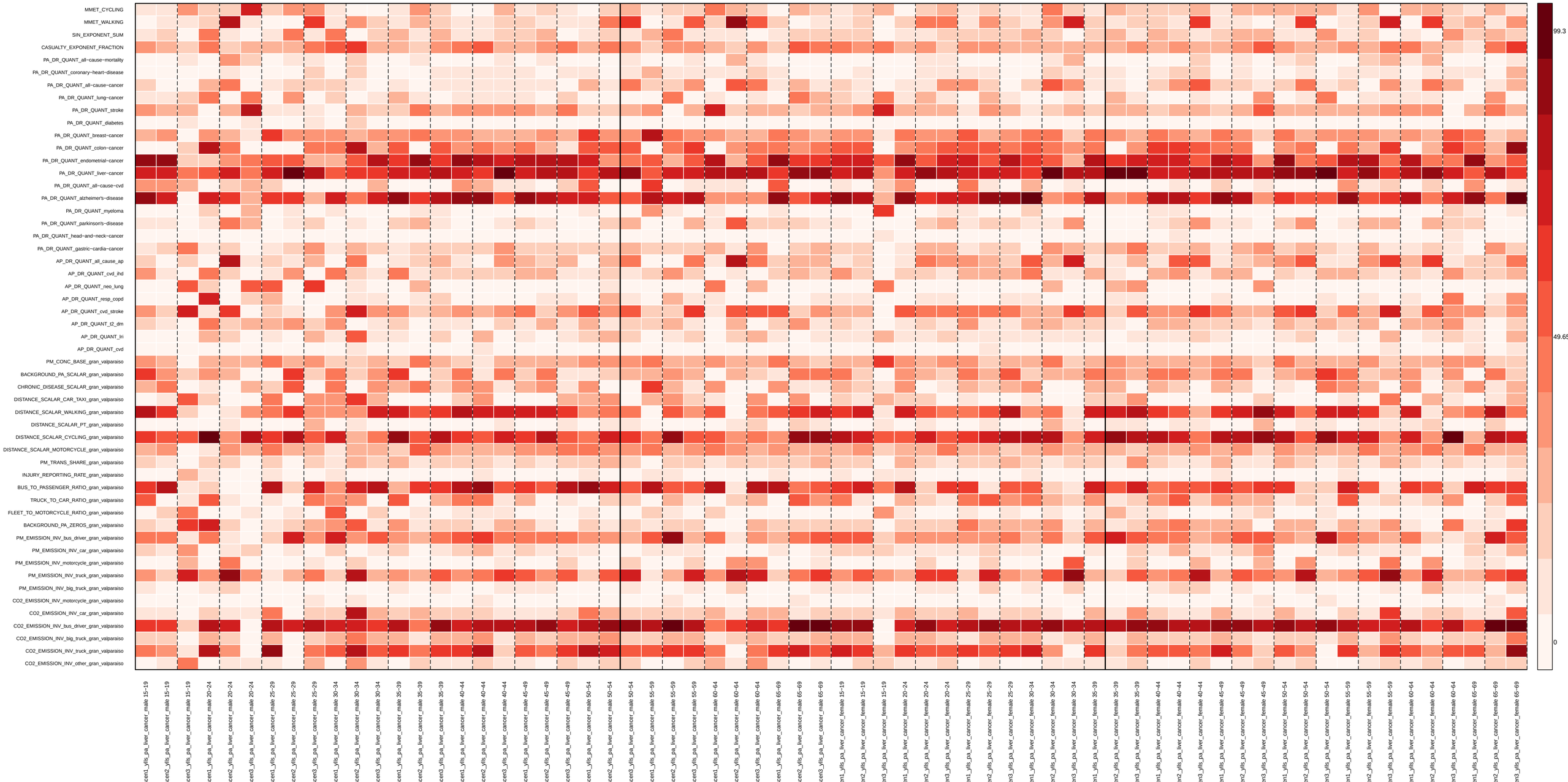
gran_valparaiso − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



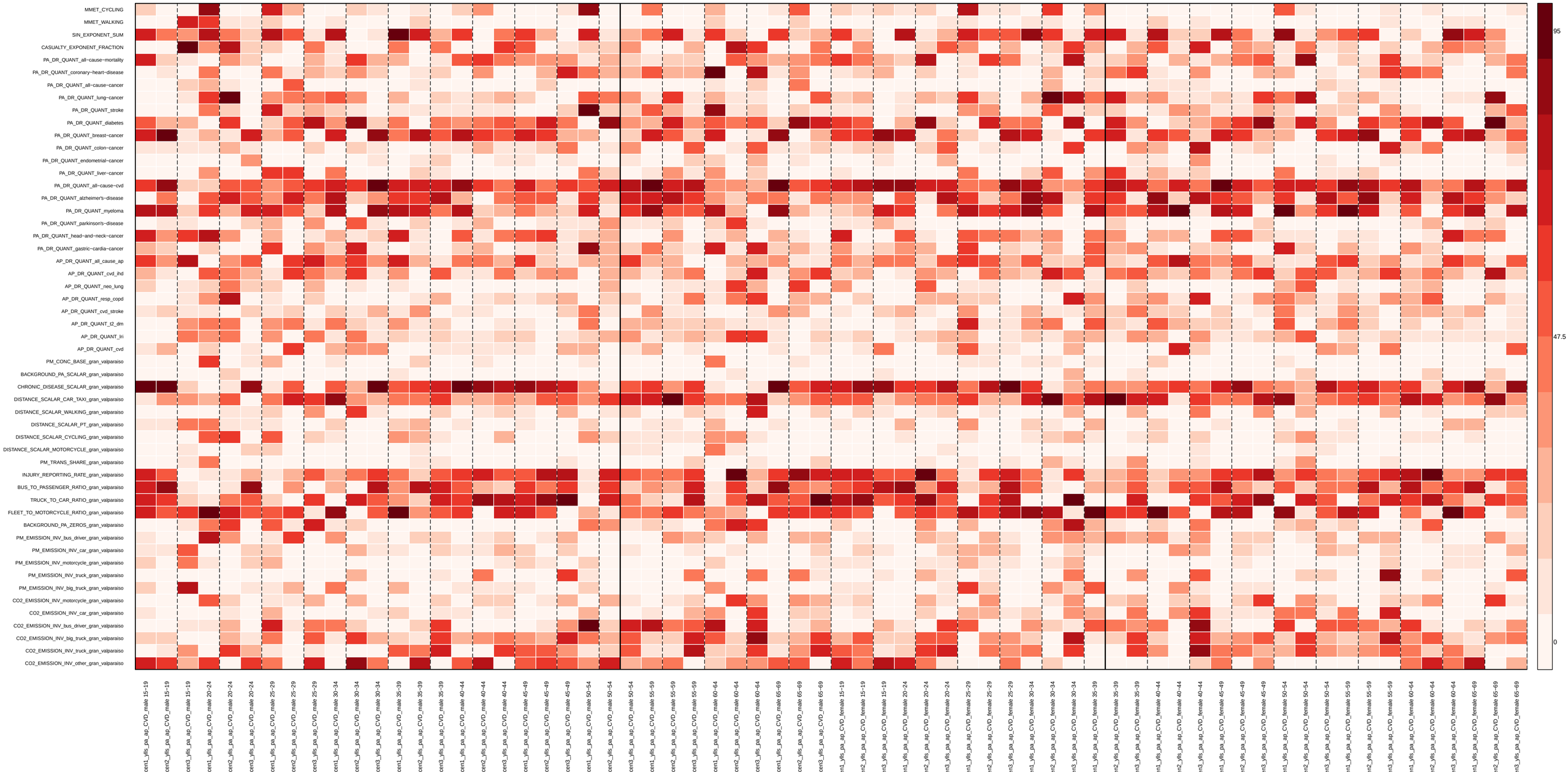
gran_valparaiso − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



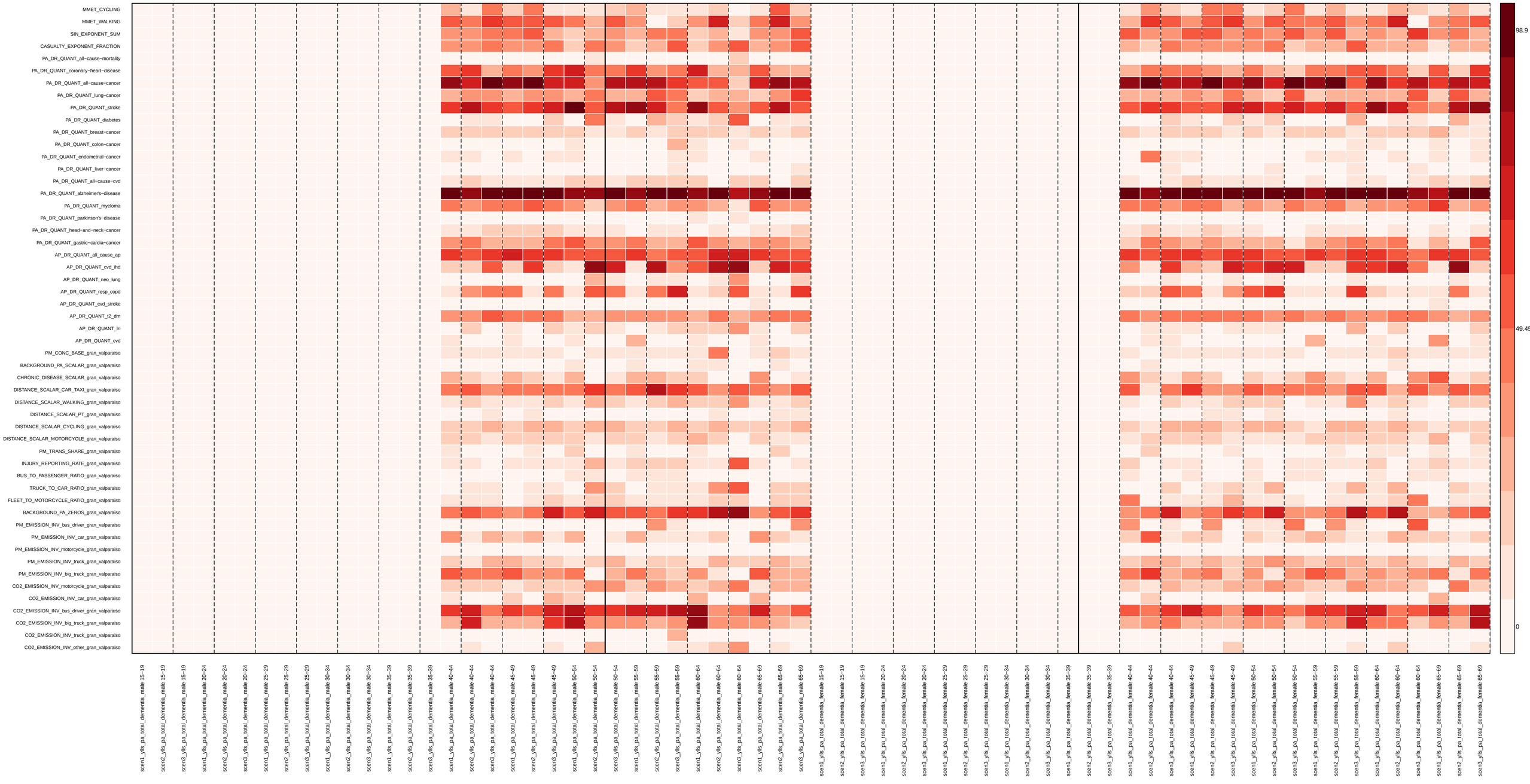
gran_valparaiso – pa_liver_cancer , No of samples: 10 –
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



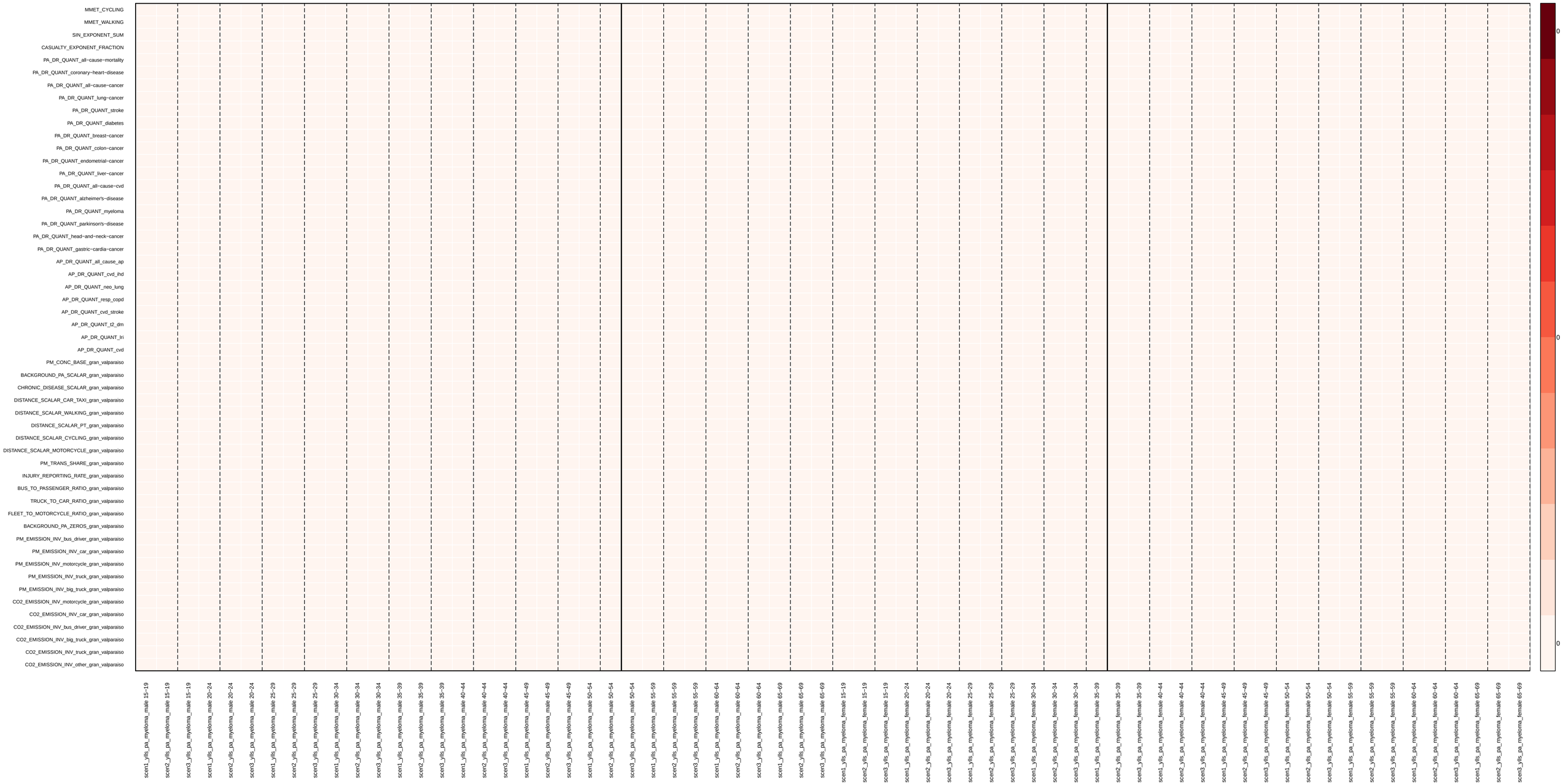
gran_valparaiso − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



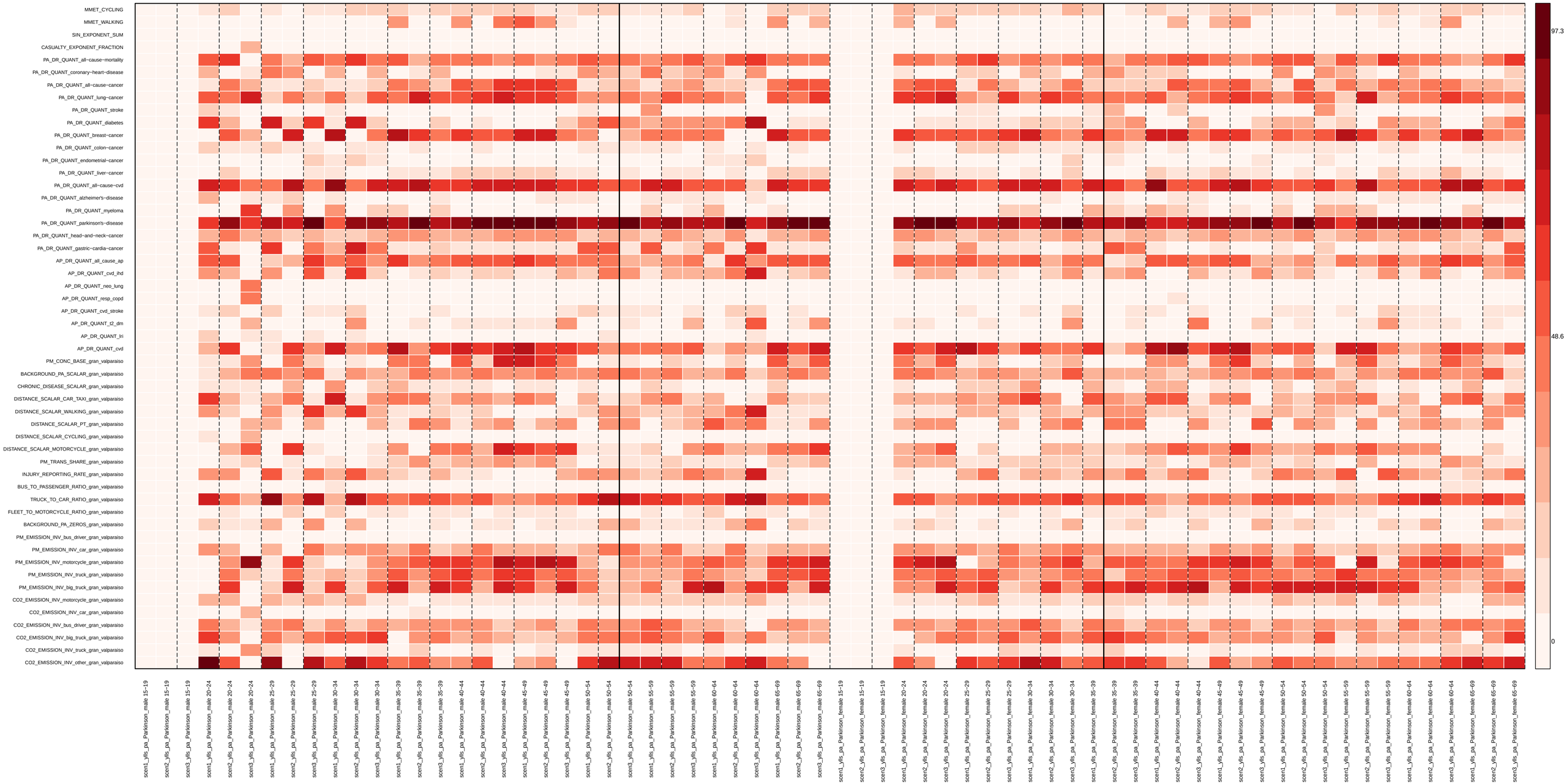
gran_valparaiso − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



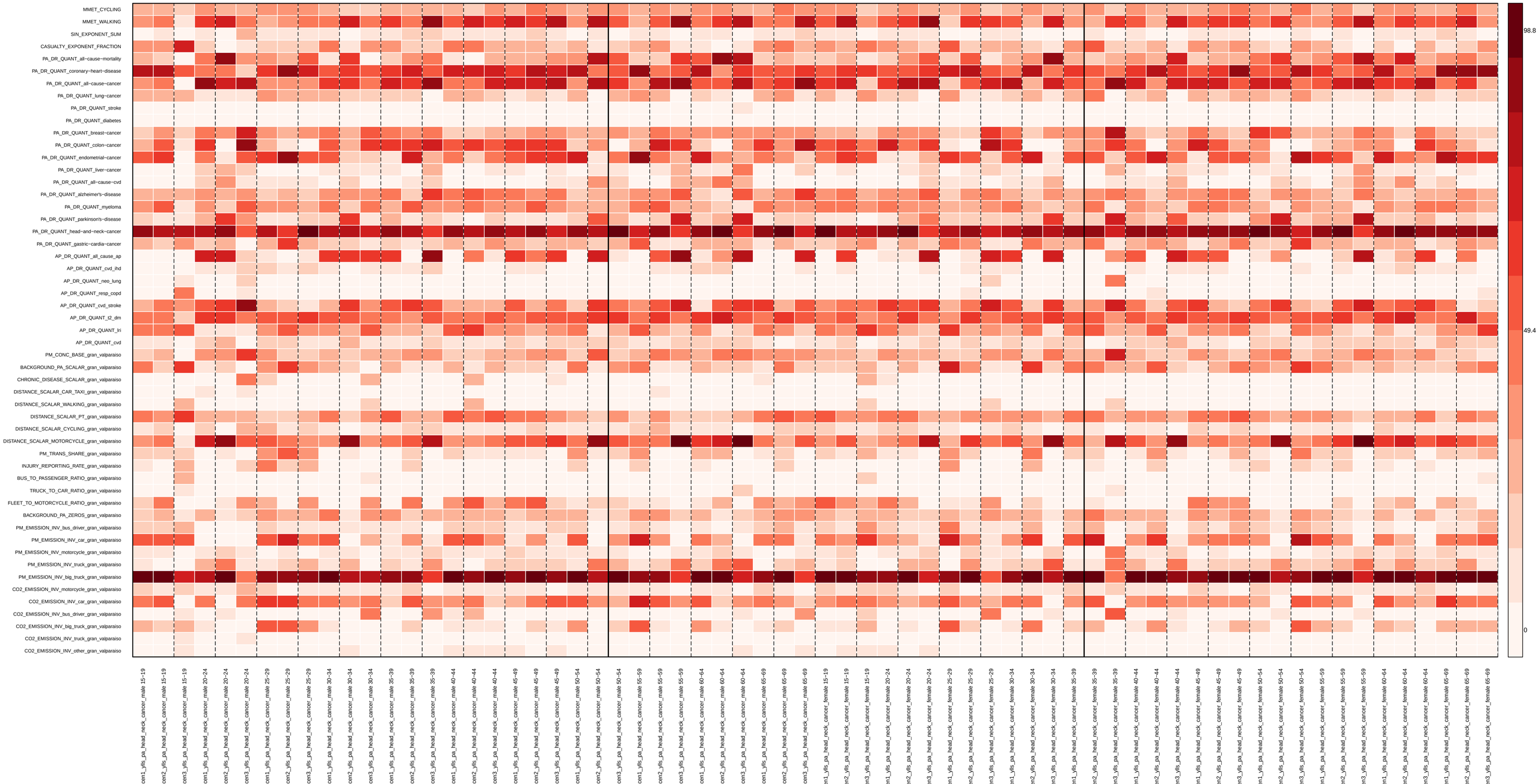
gran_valparaiso − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



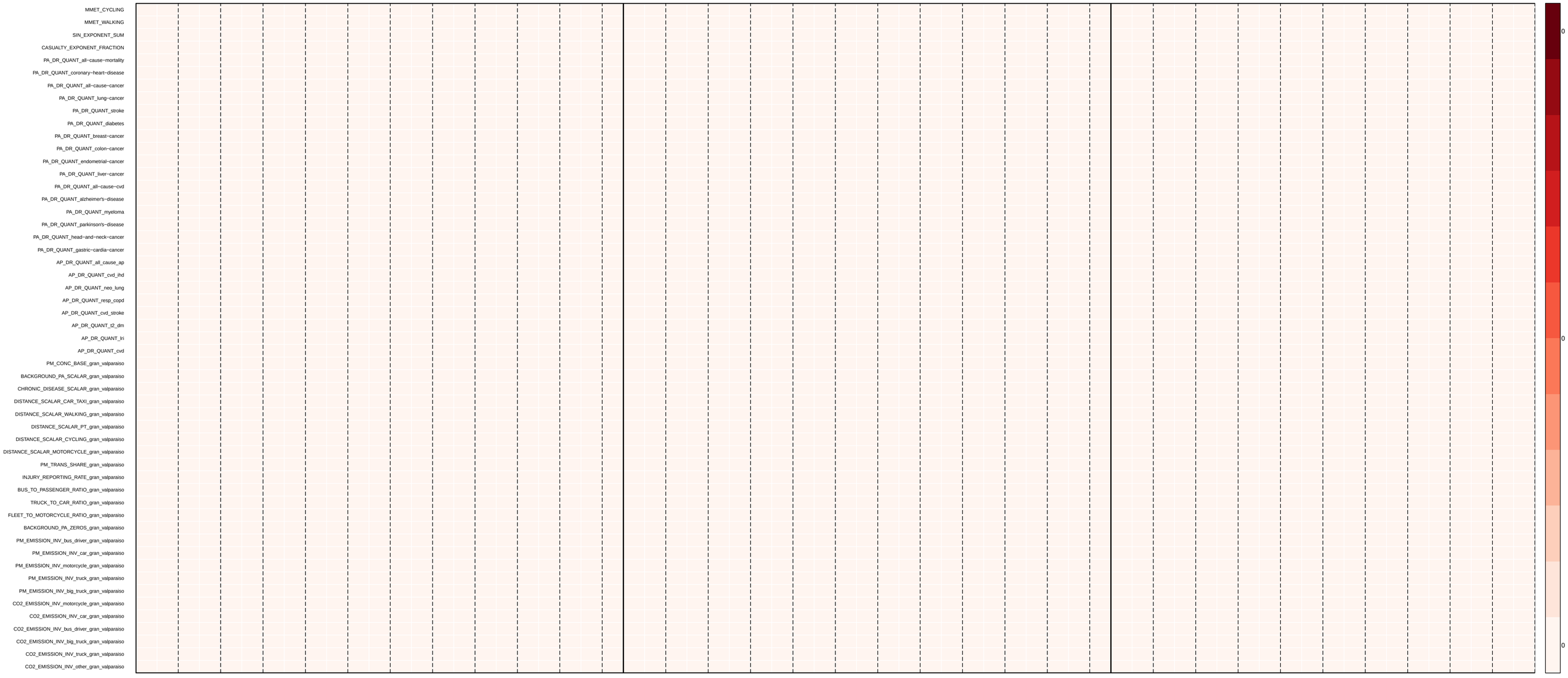
gran_valparaiso − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



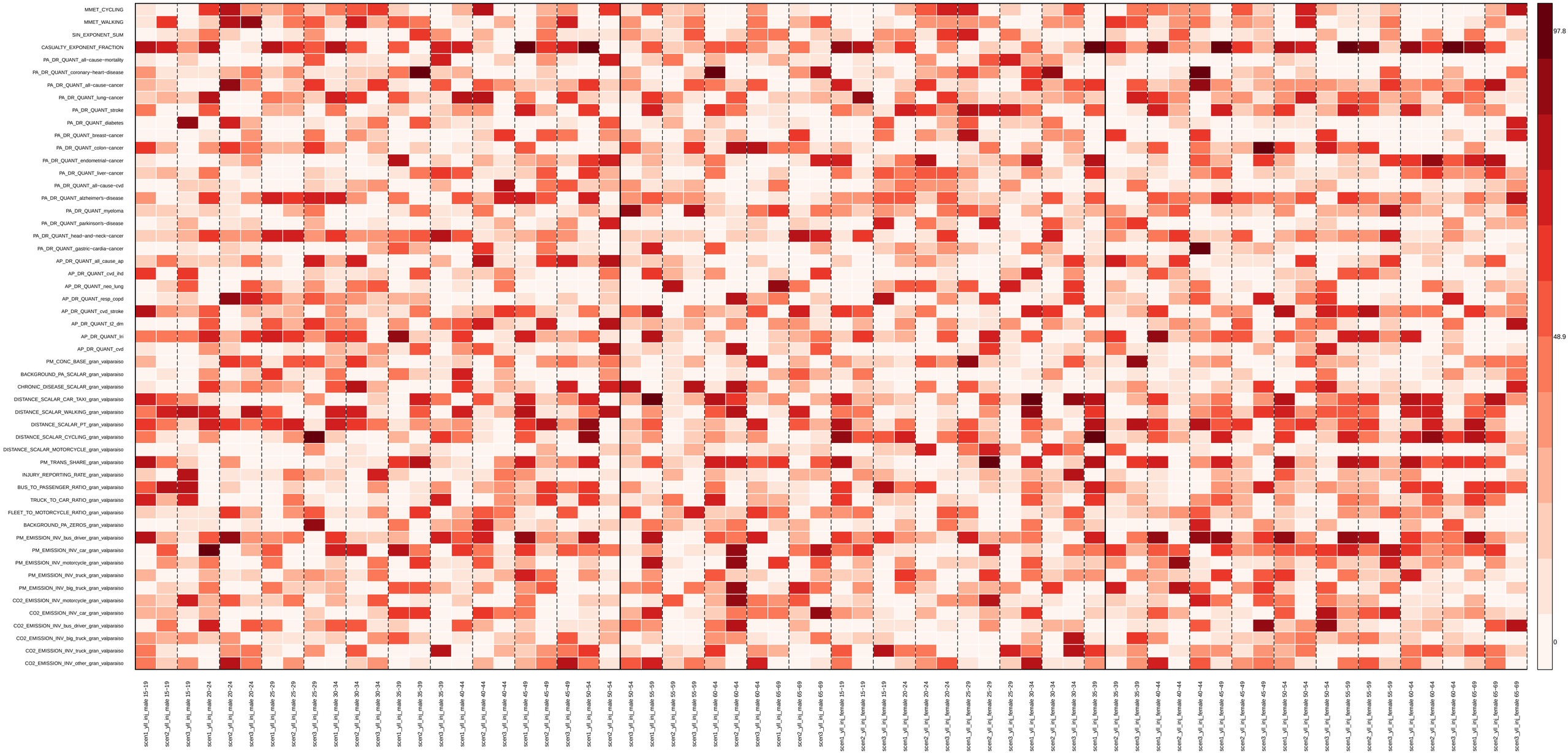
gran_valparaiso − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



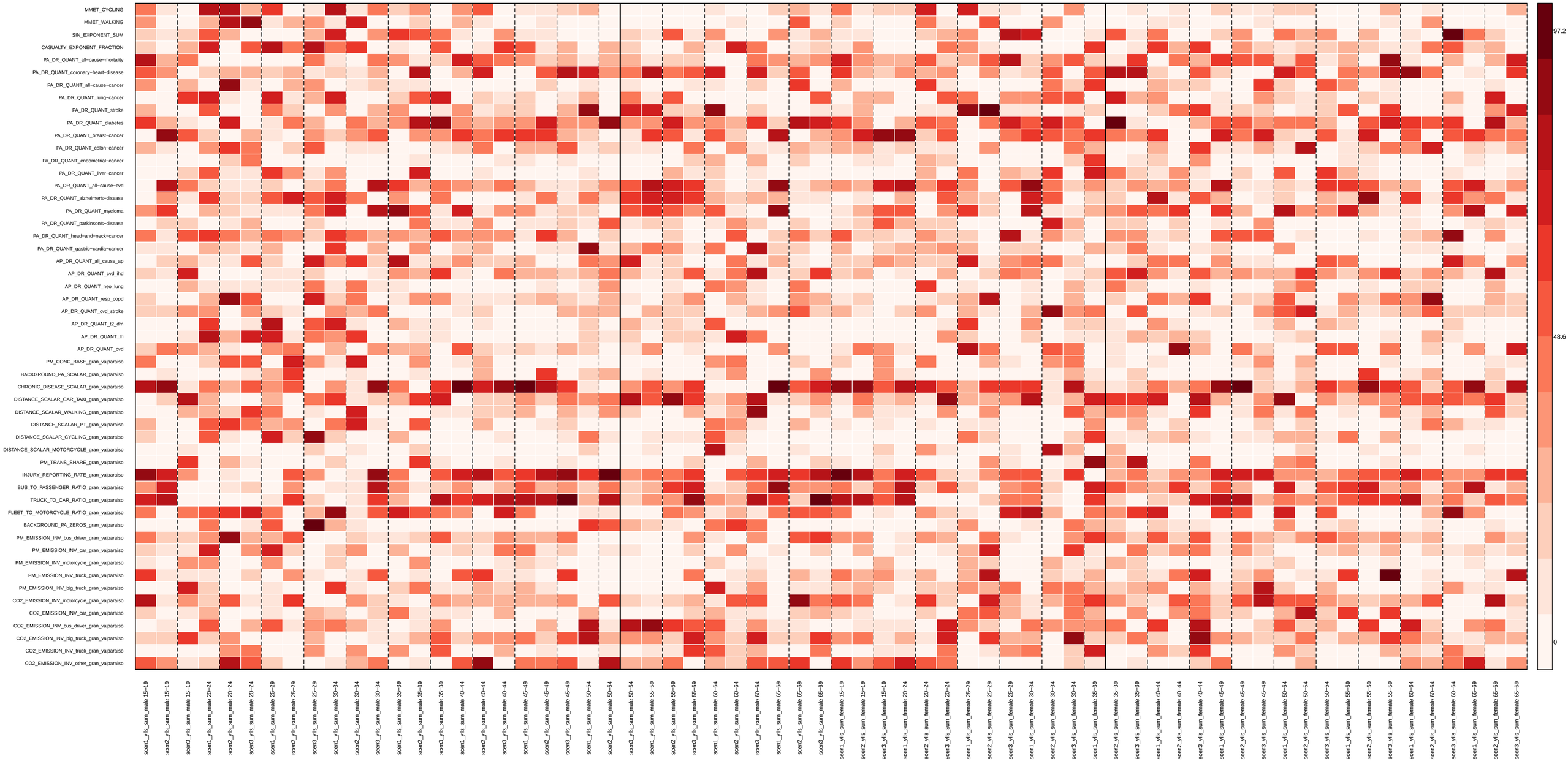
gran_valparaiso − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



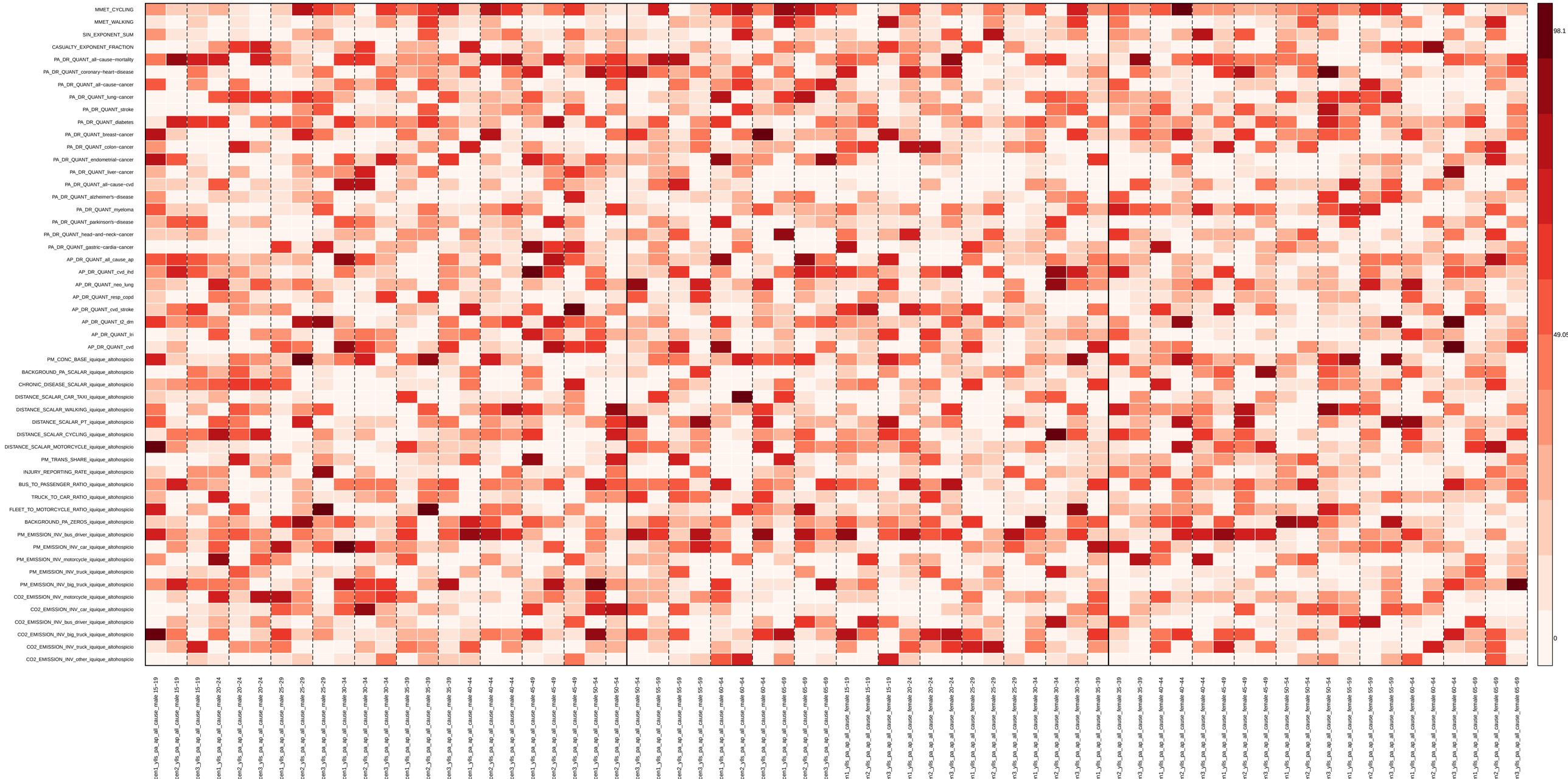
gran_valparaiso − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



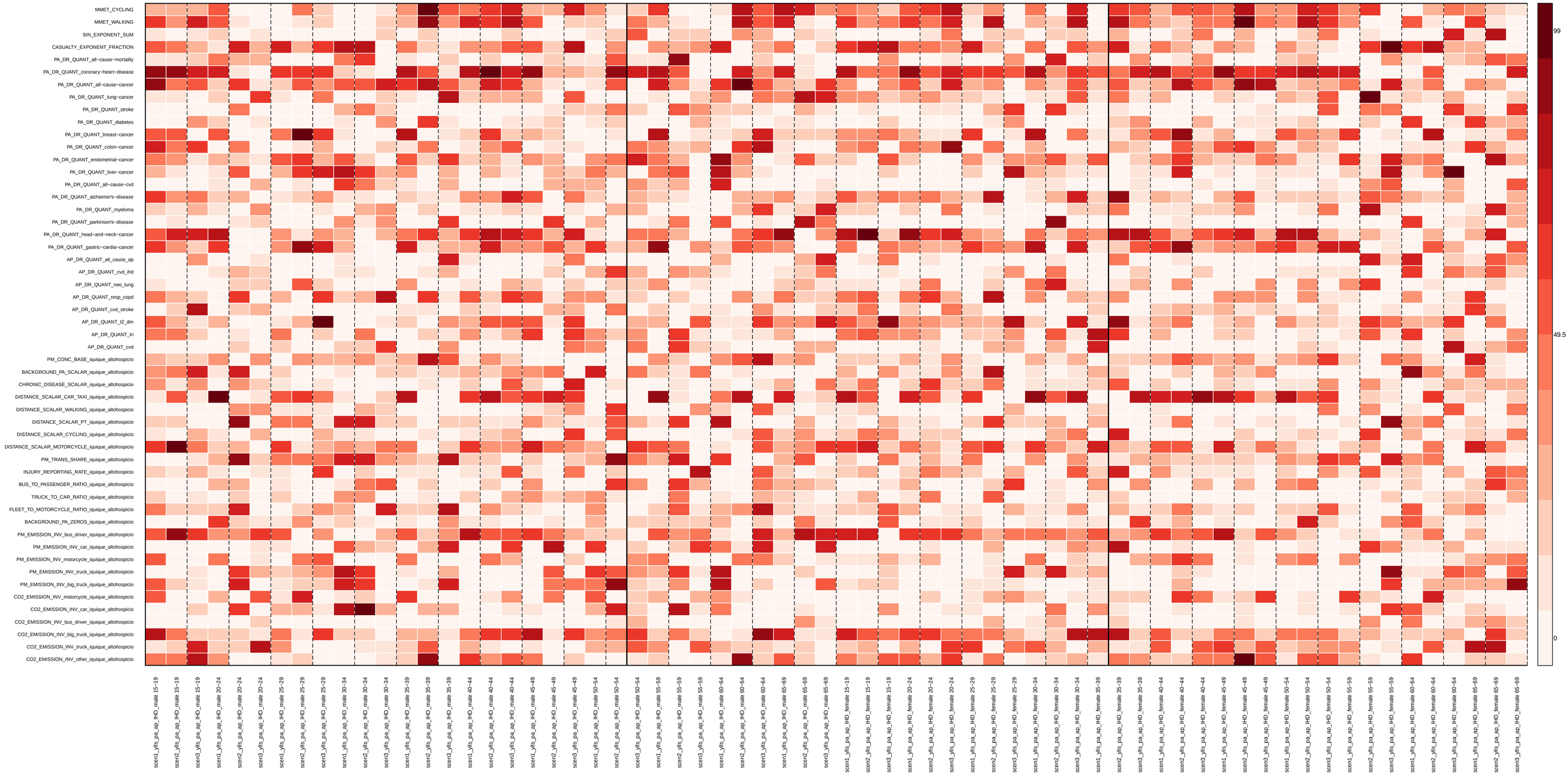
gran_valparaiso − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



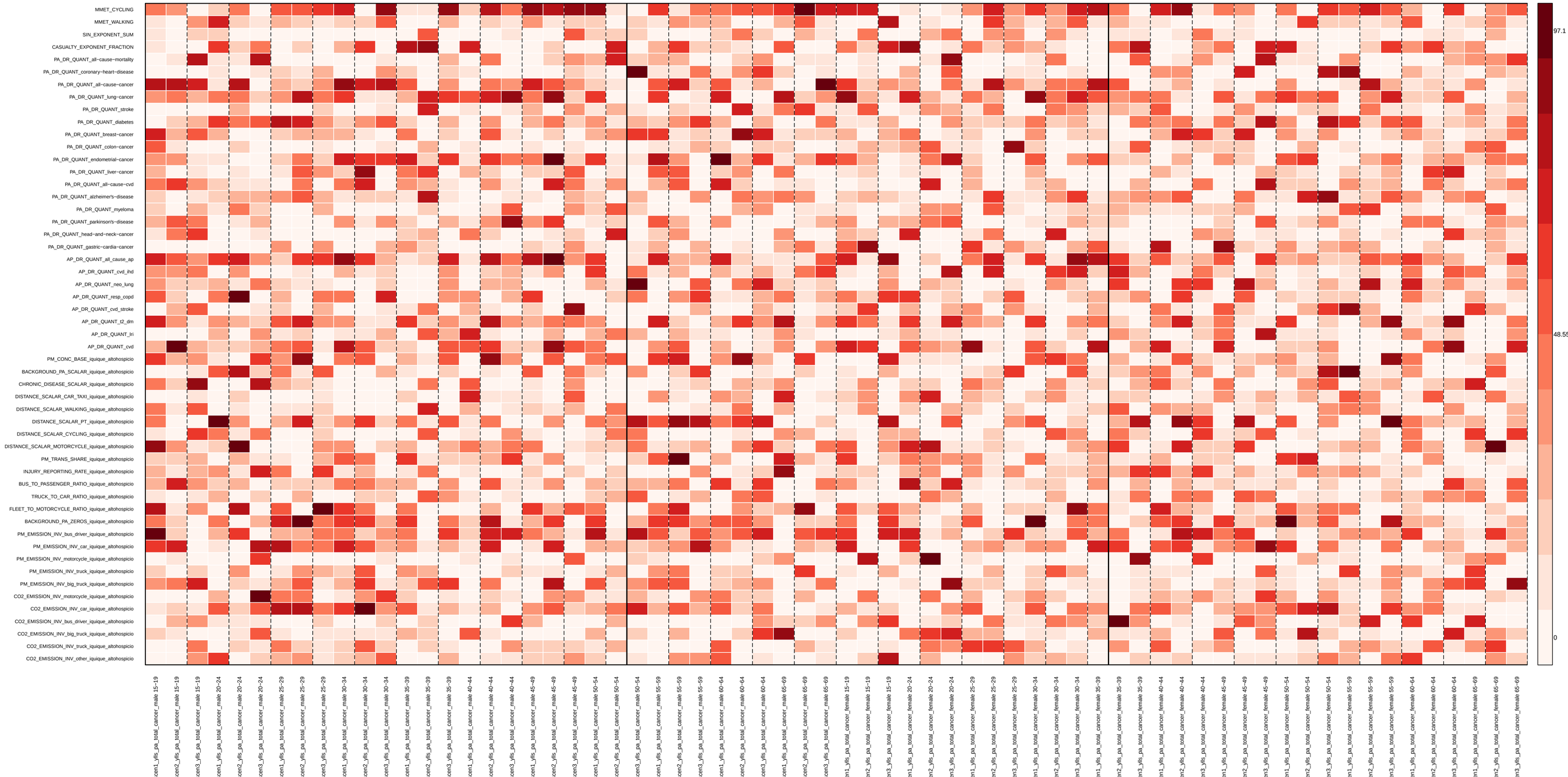
iquique_altohospicio − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



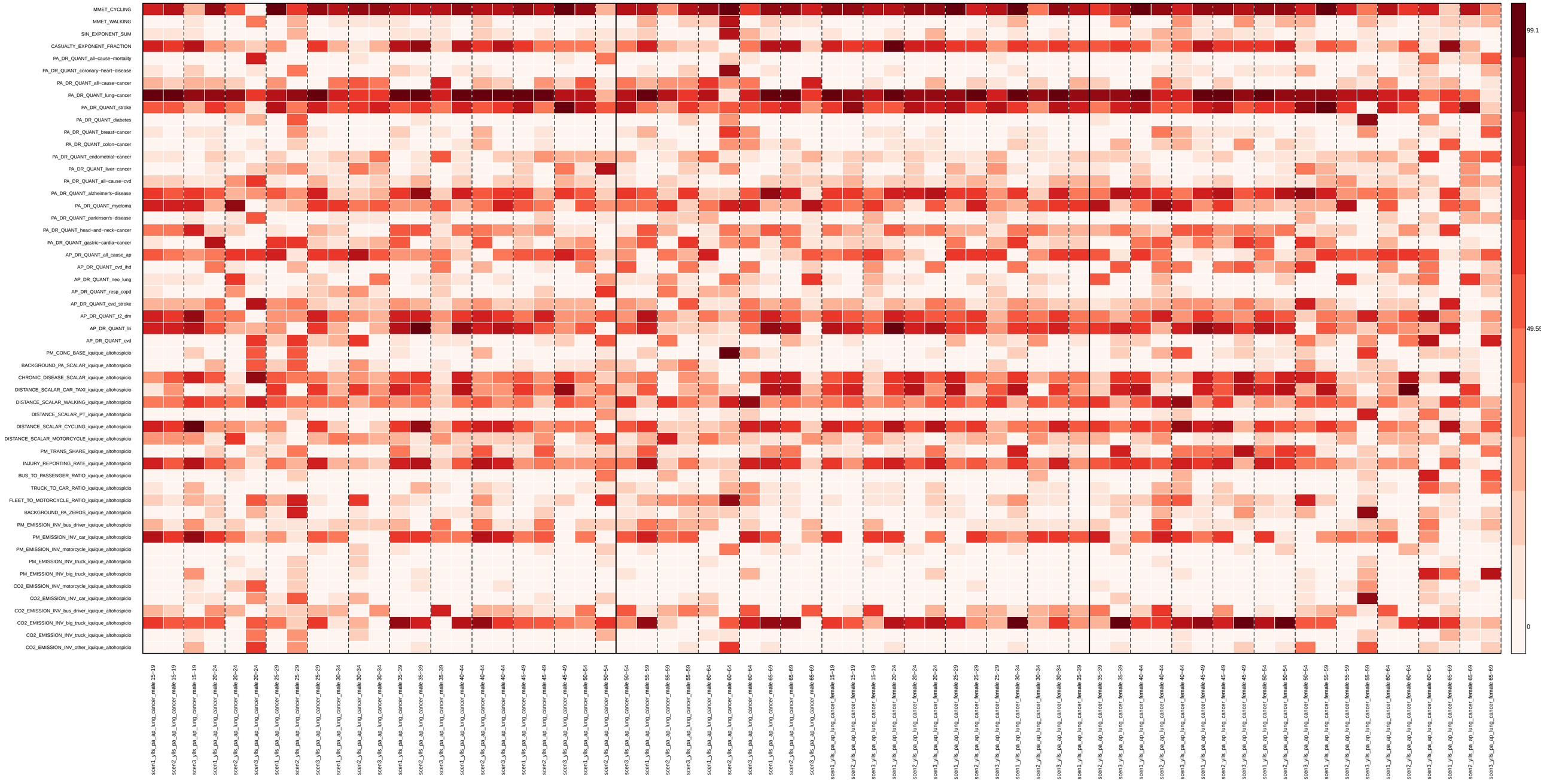
iquique_altohospicio − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



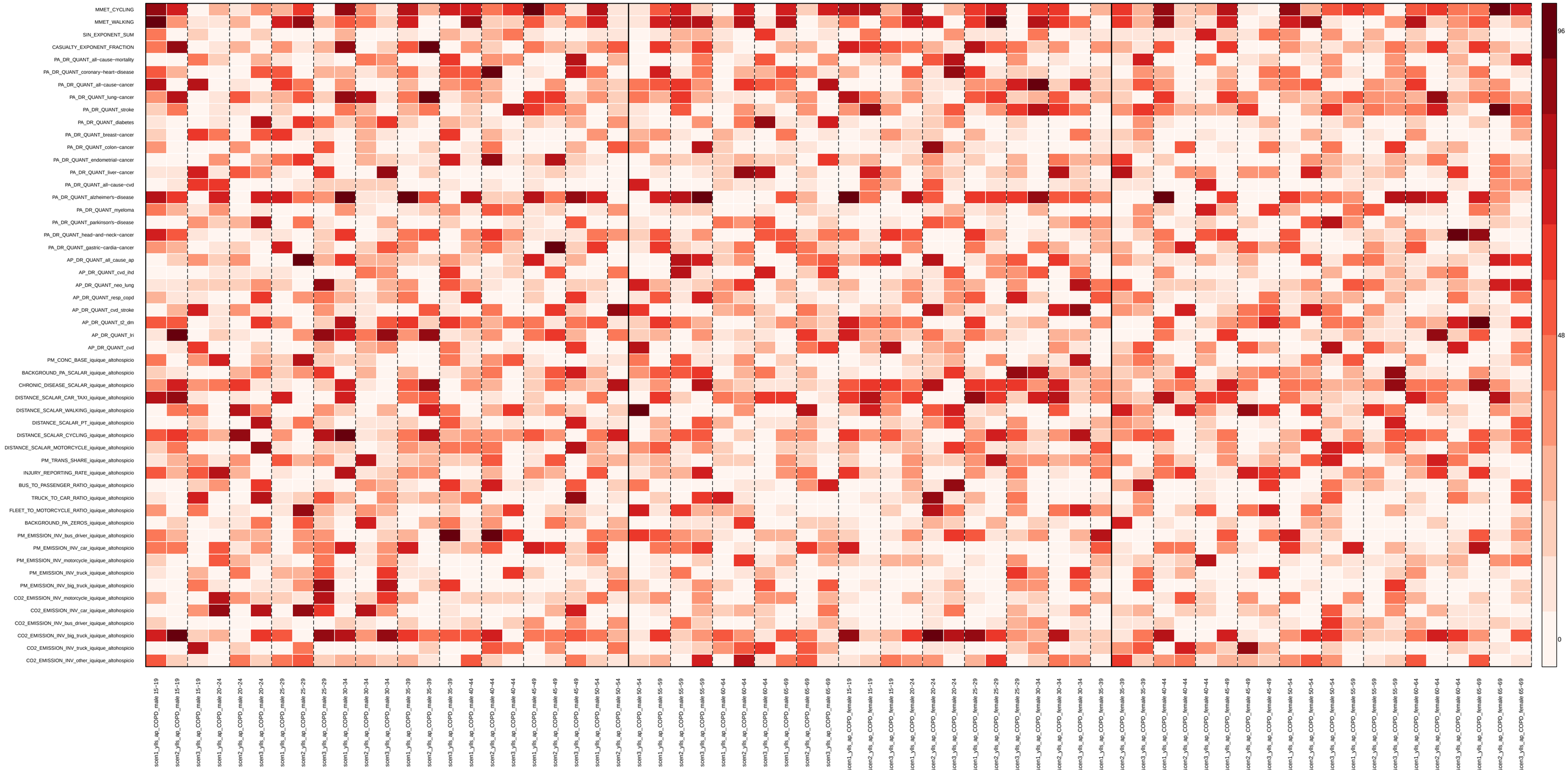
iquique_altohospicio – pa_total_cancer , No of samples: 10 –
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



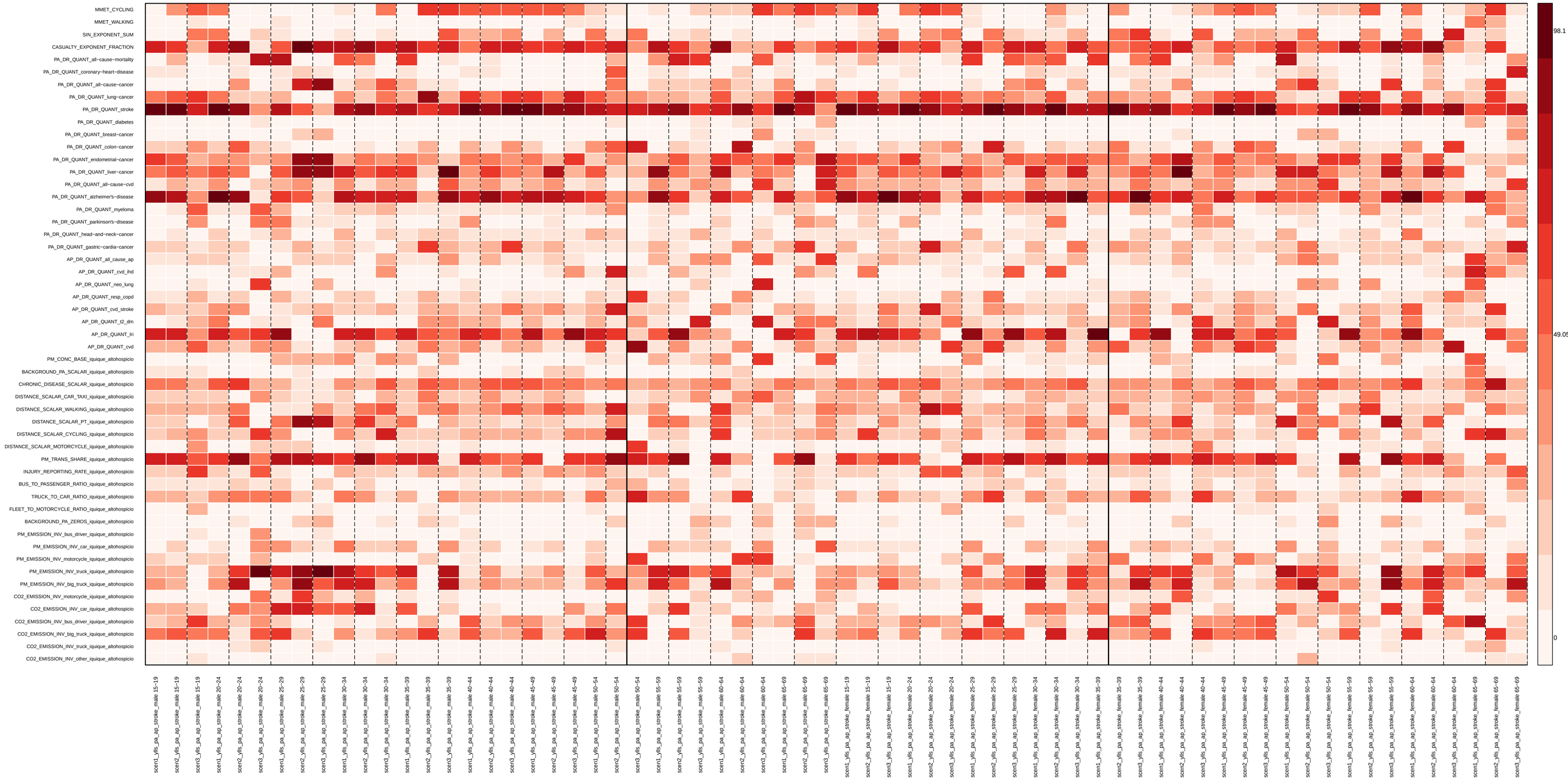
iquique_altohospicio – pa_ap_lung_cancer , No of samples: 10 –
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



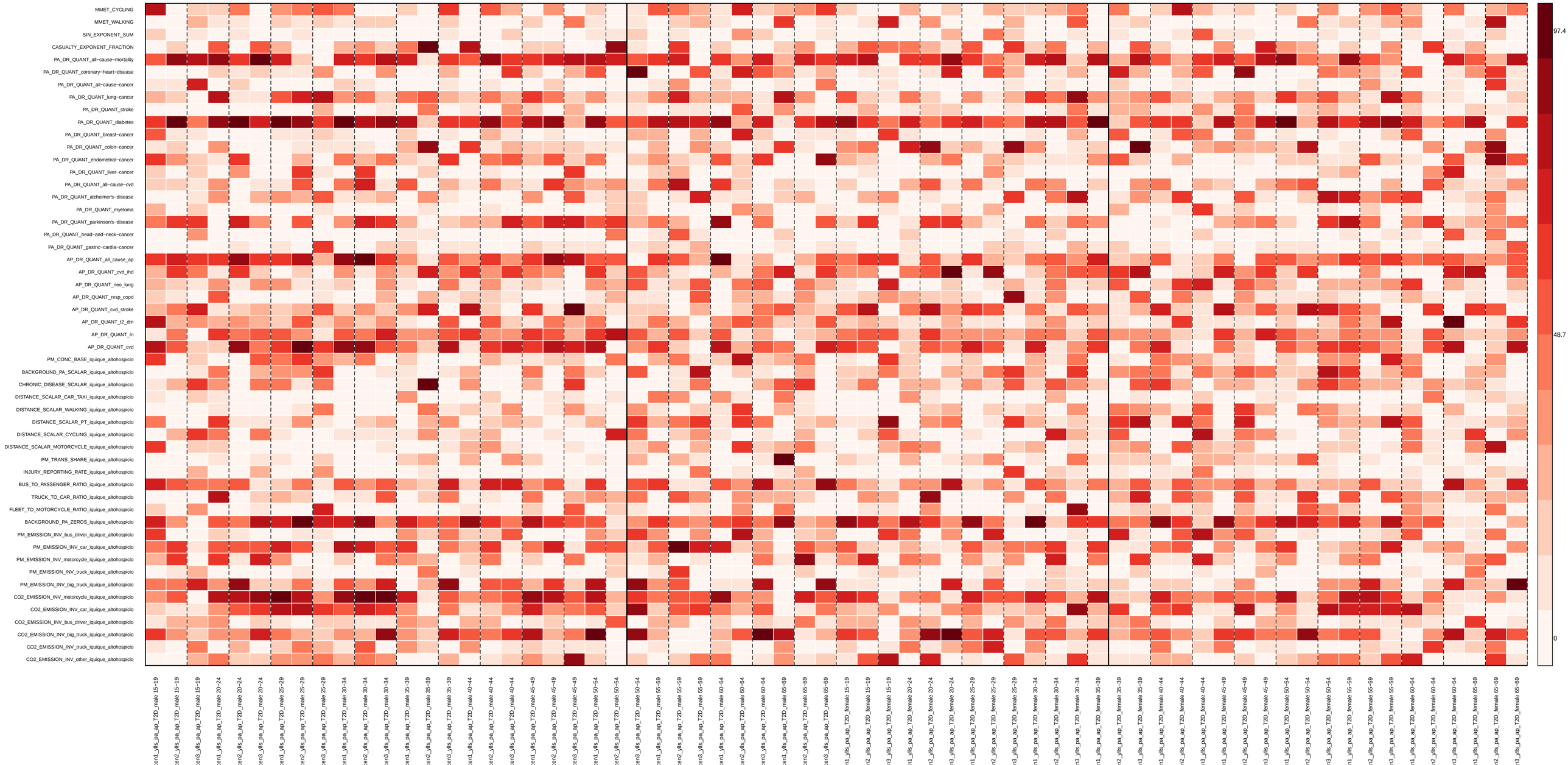
iquique_altohospicio − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



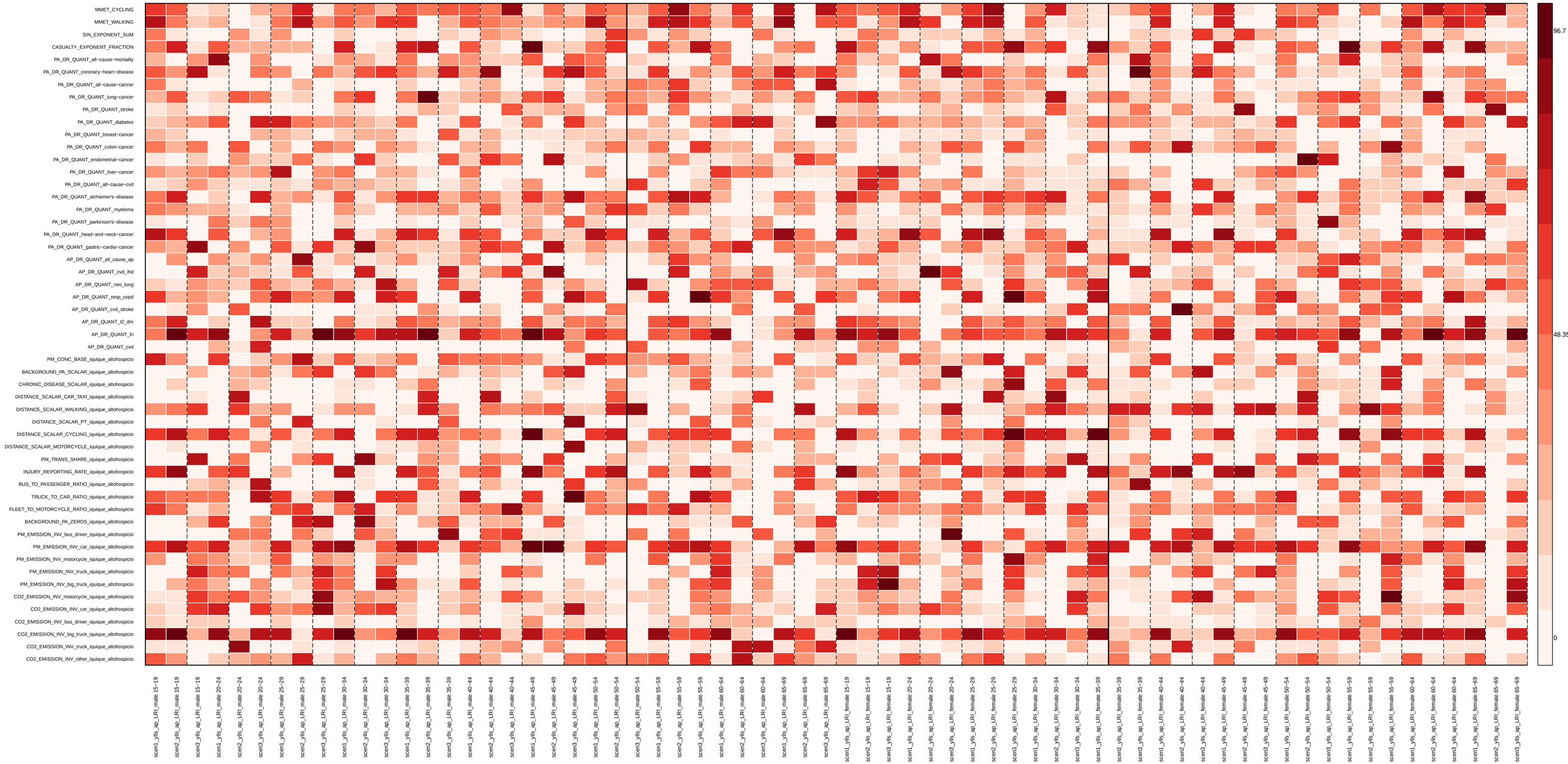
iquique_altohospicio − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



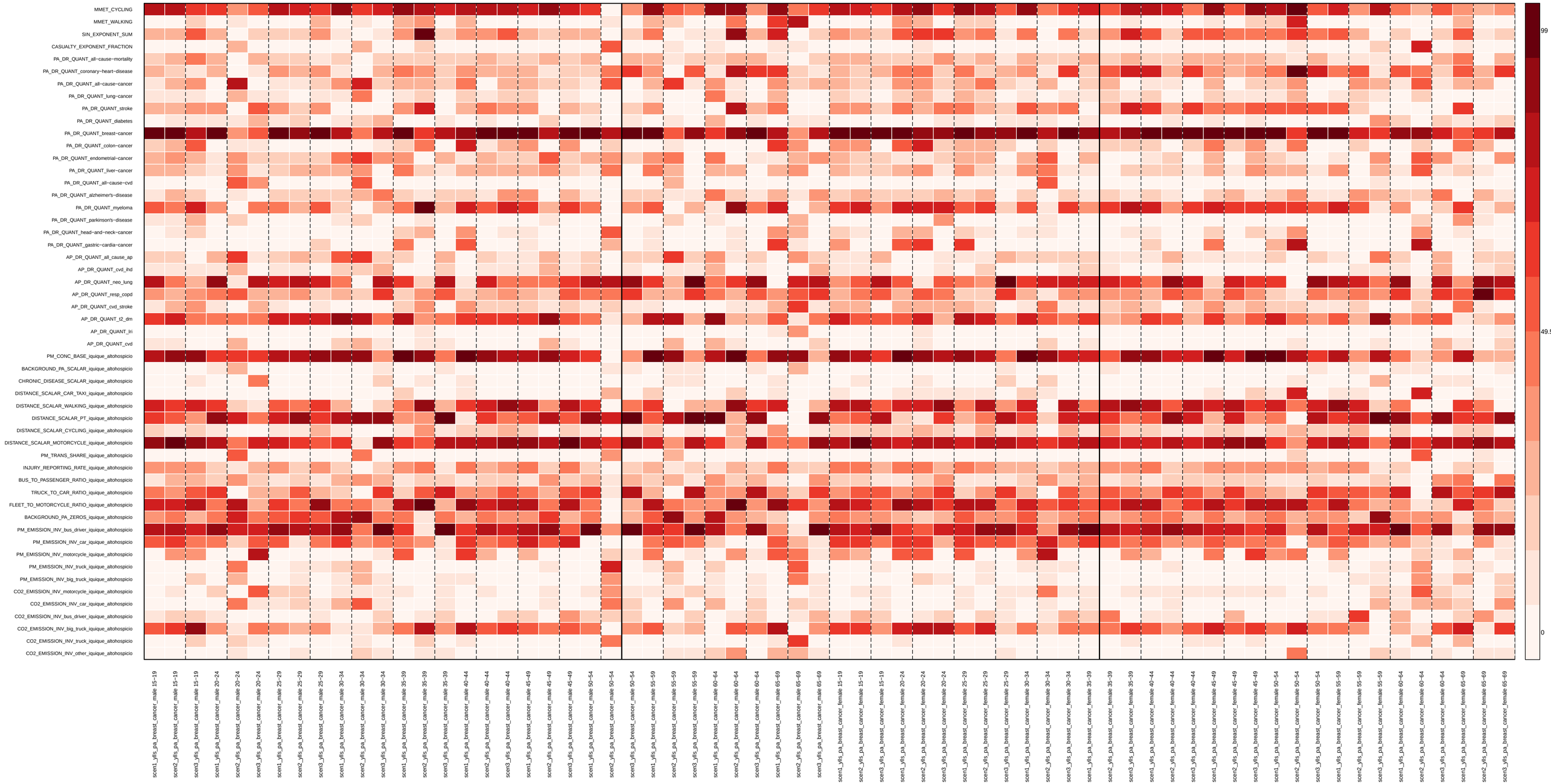
iquique_altohospicio − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



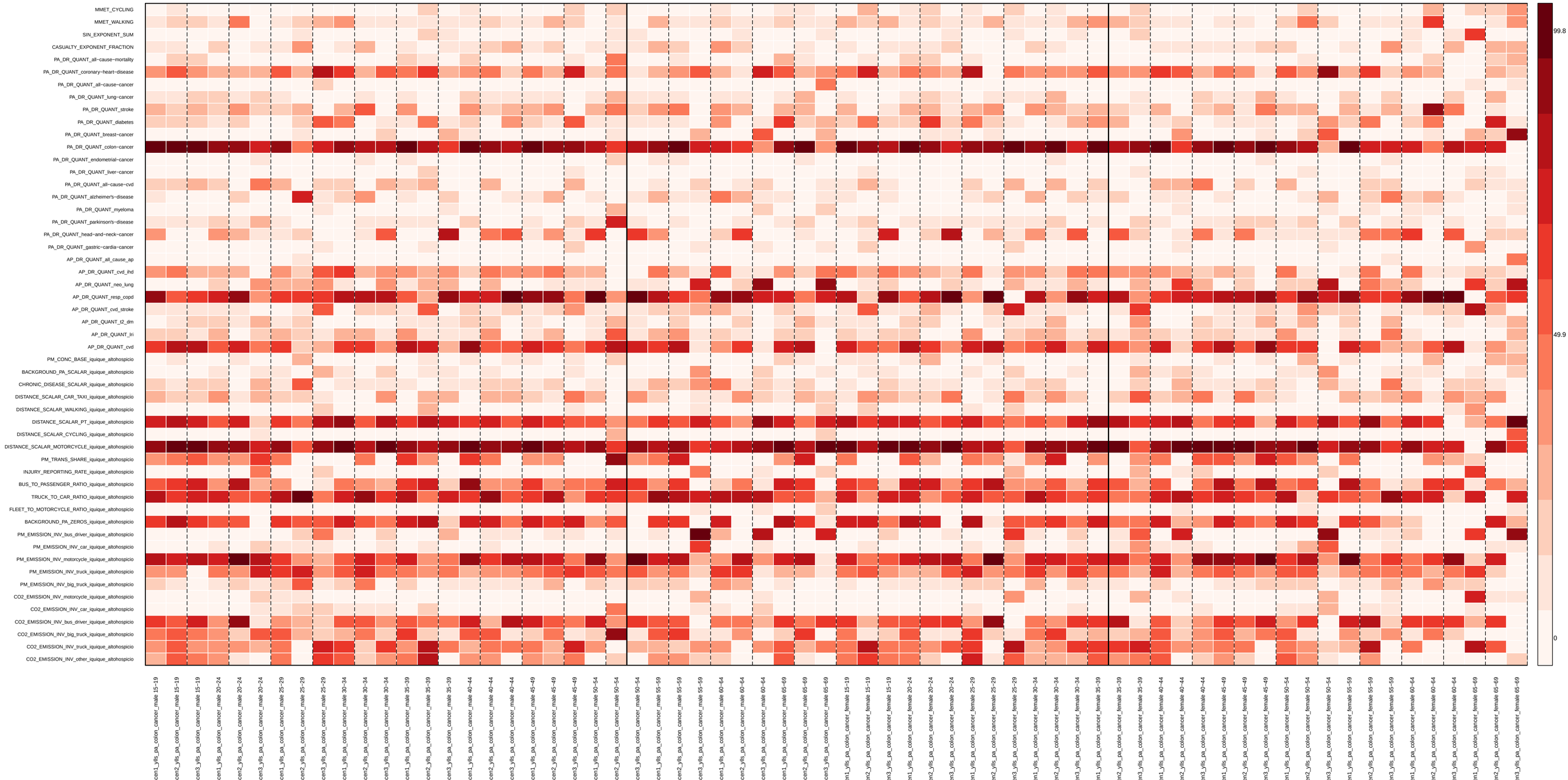
iquique_altohospicio − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



iquique_altohospicio − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

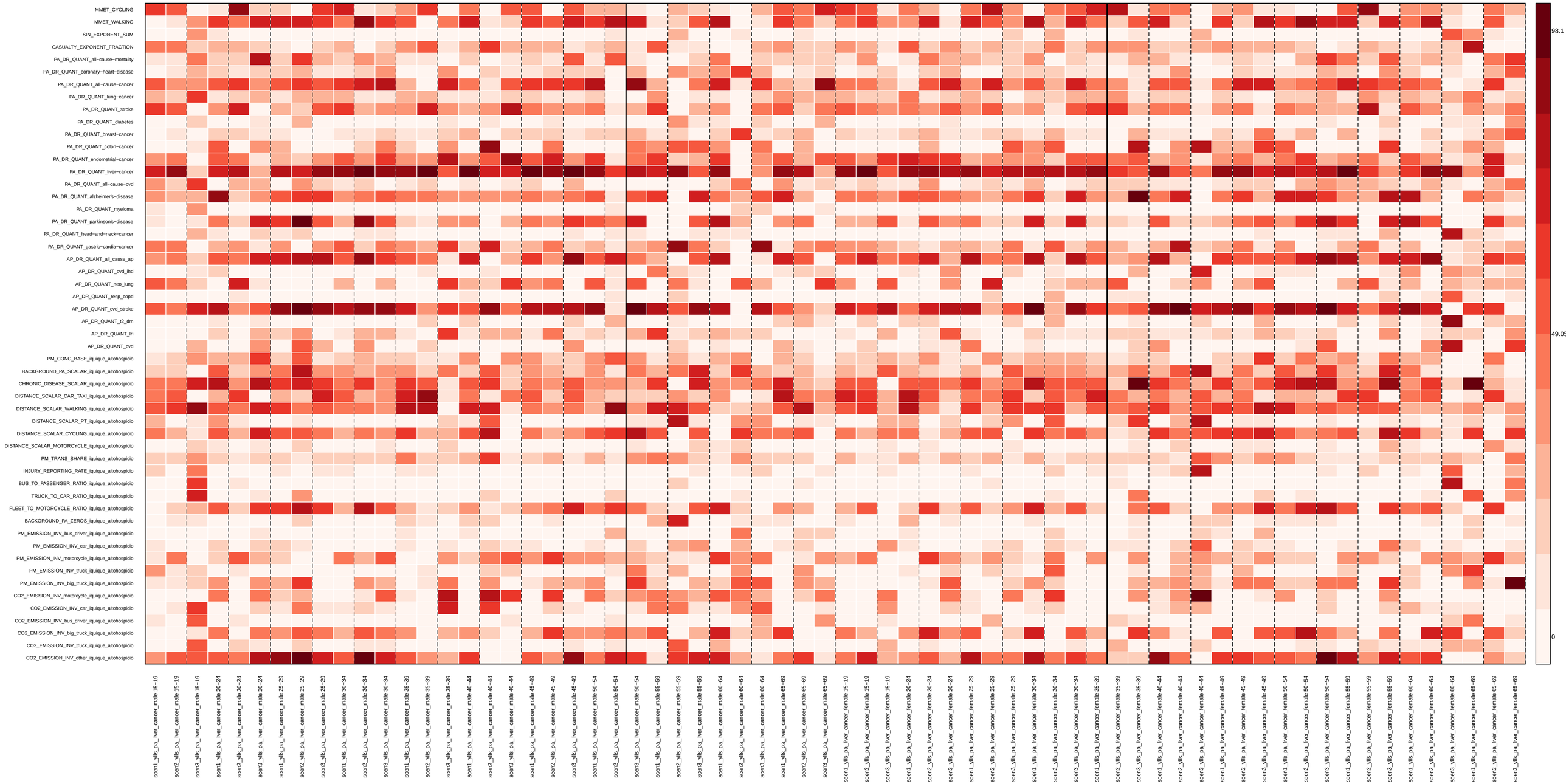


iquique_altohospicio − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

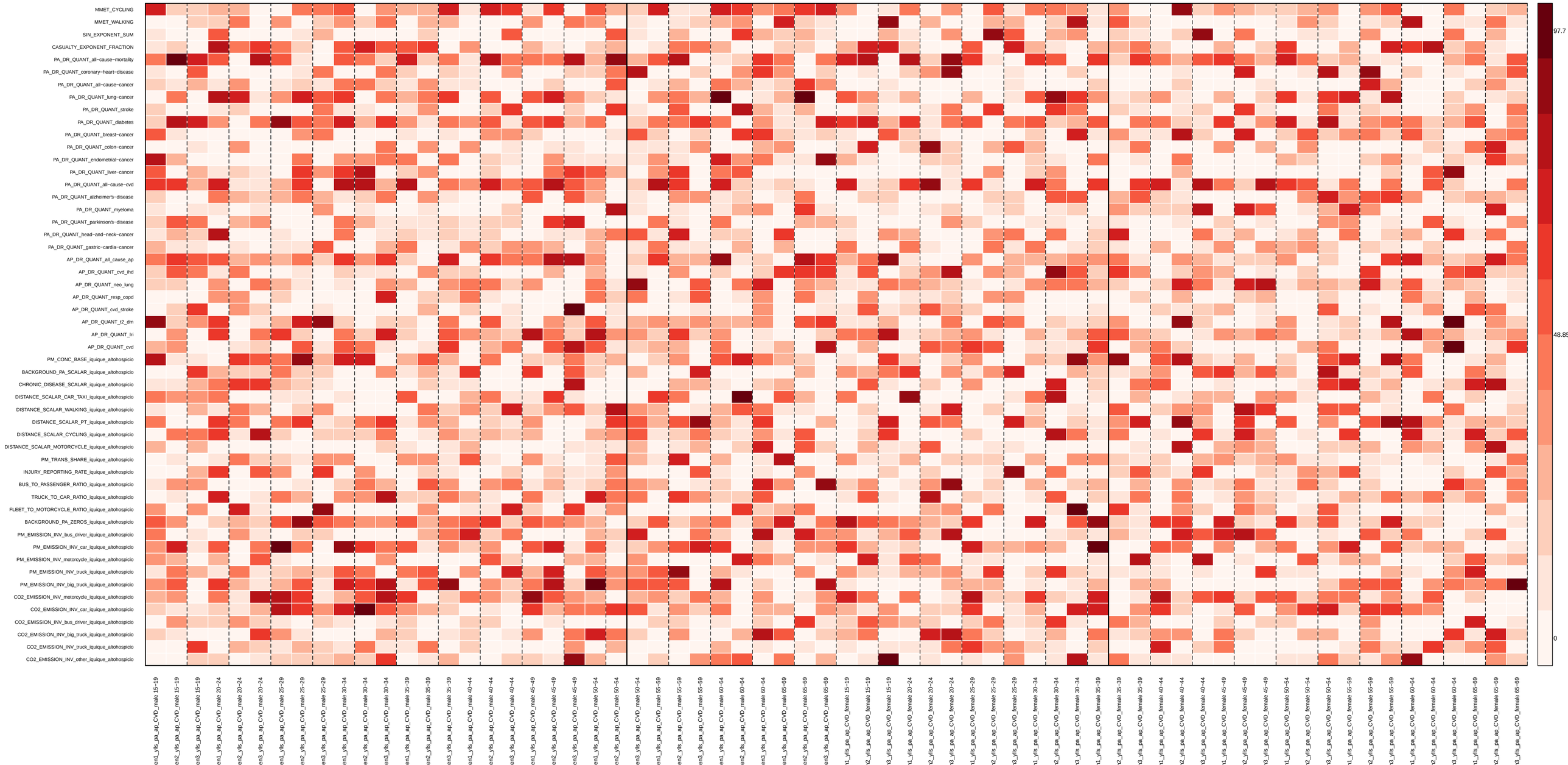


iquique_altohospicio – pa_endo_cancer , No of samples: 10 –
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

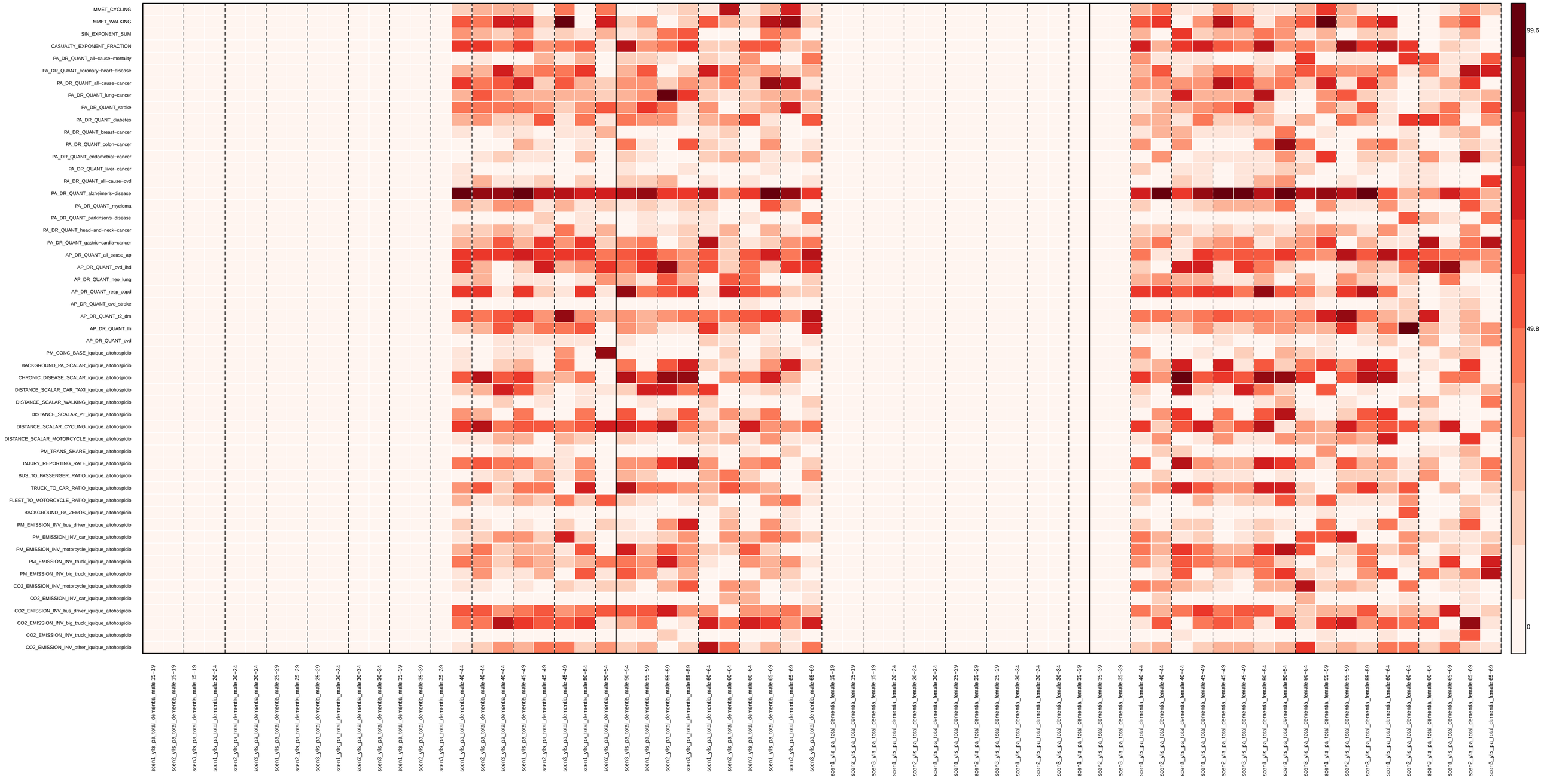
iquique_altohospicio − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



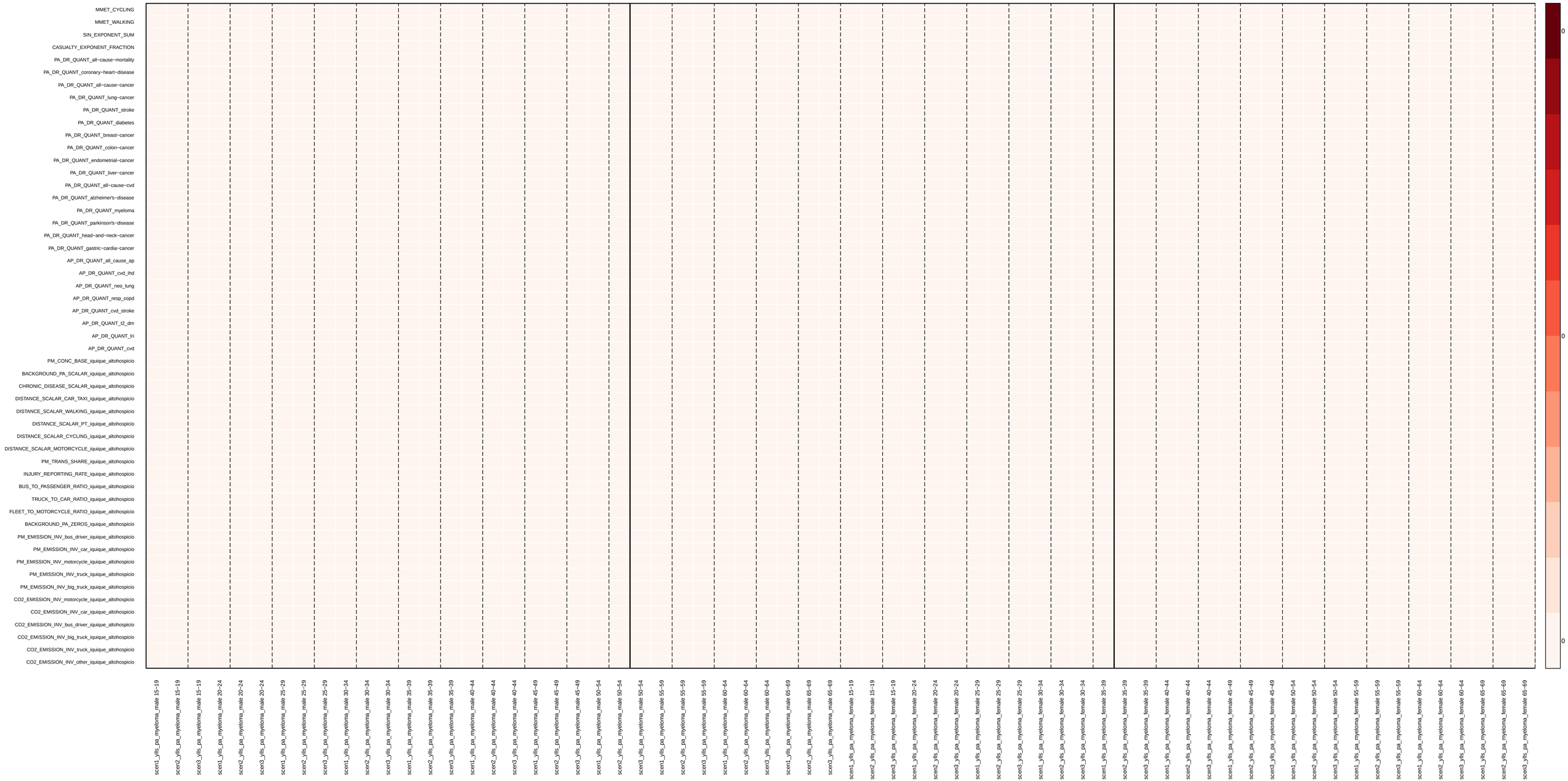
iquique_altohospicio − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



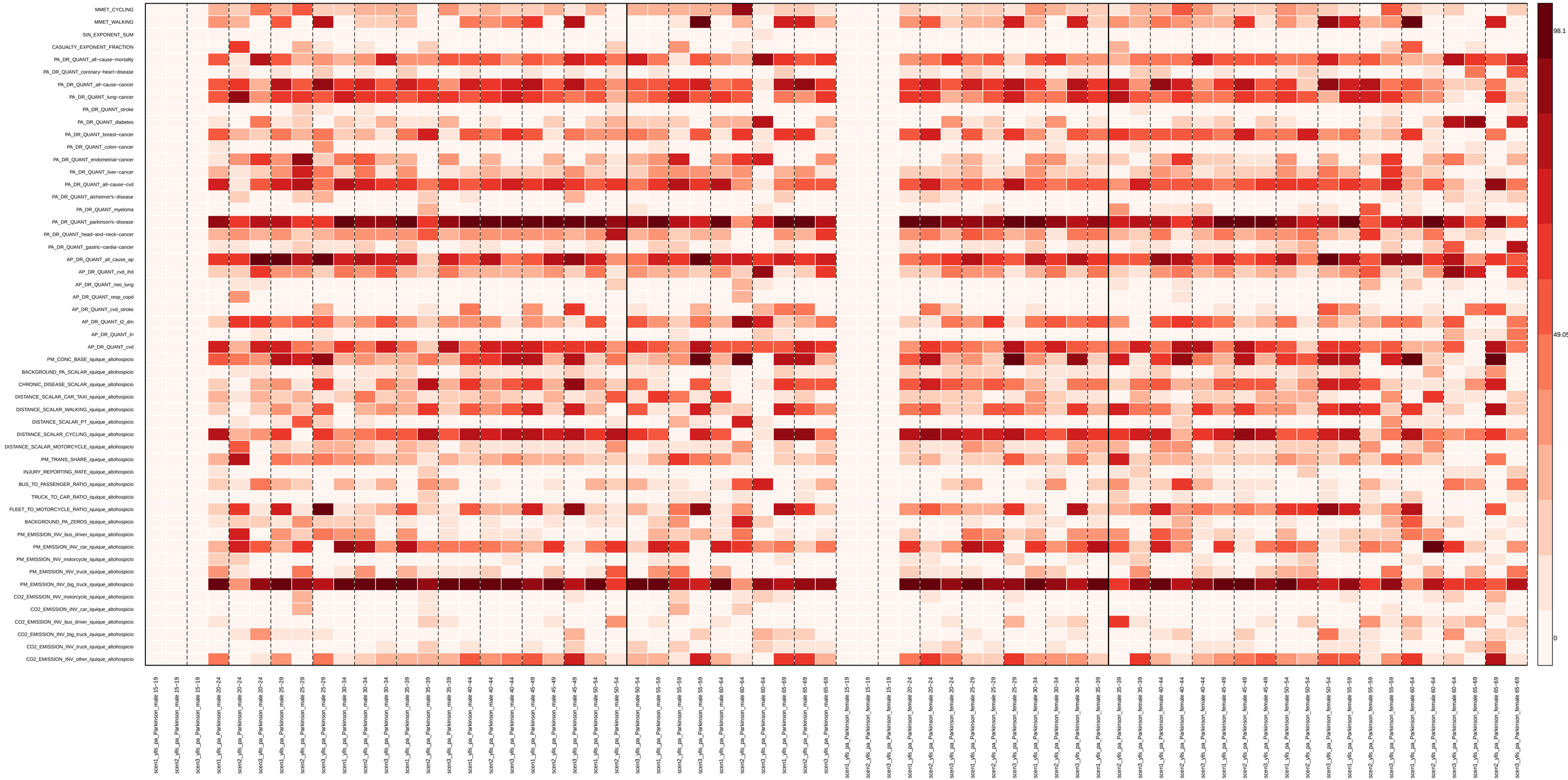
iquique_altohospicio − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



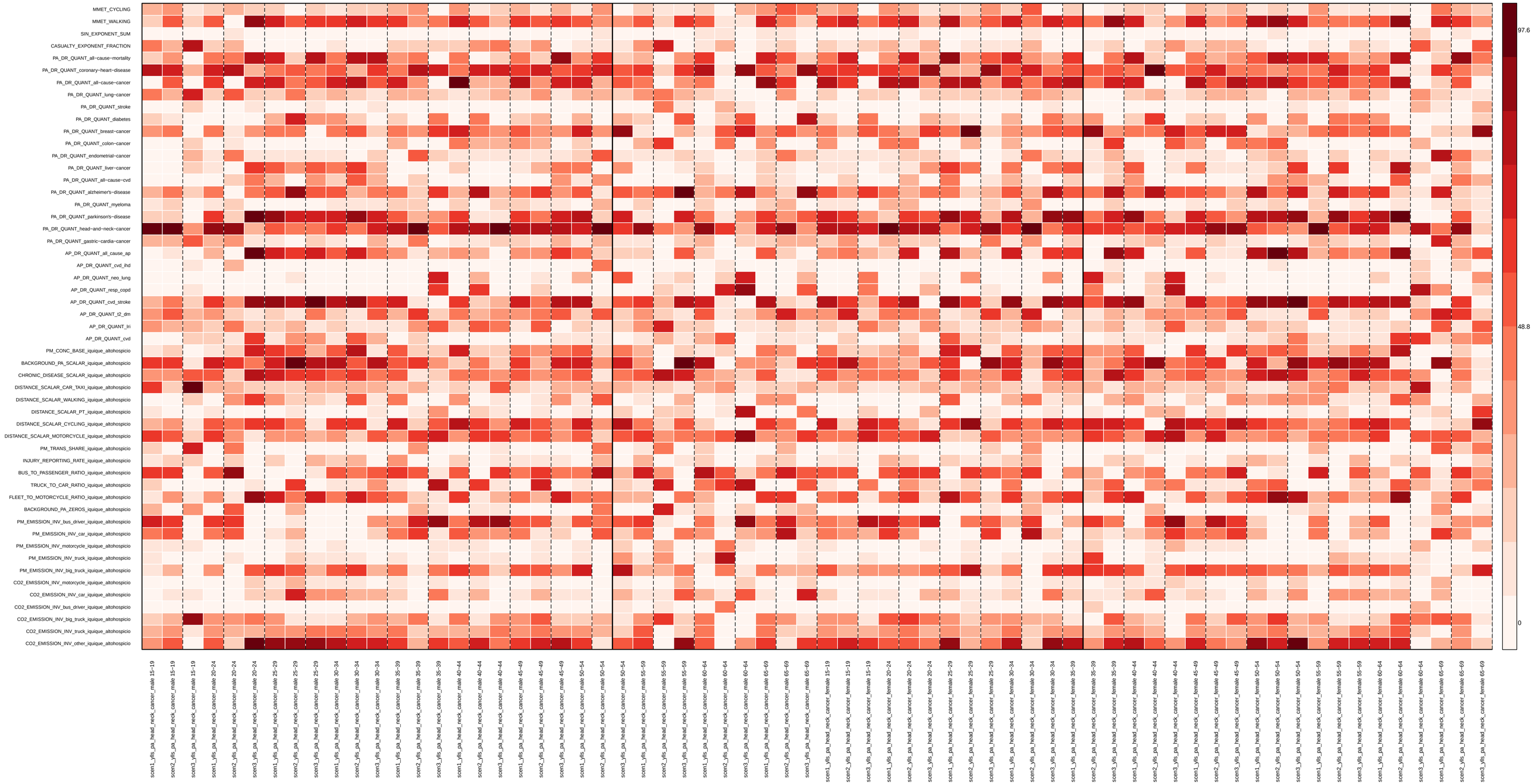
iquique_altohospicio − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



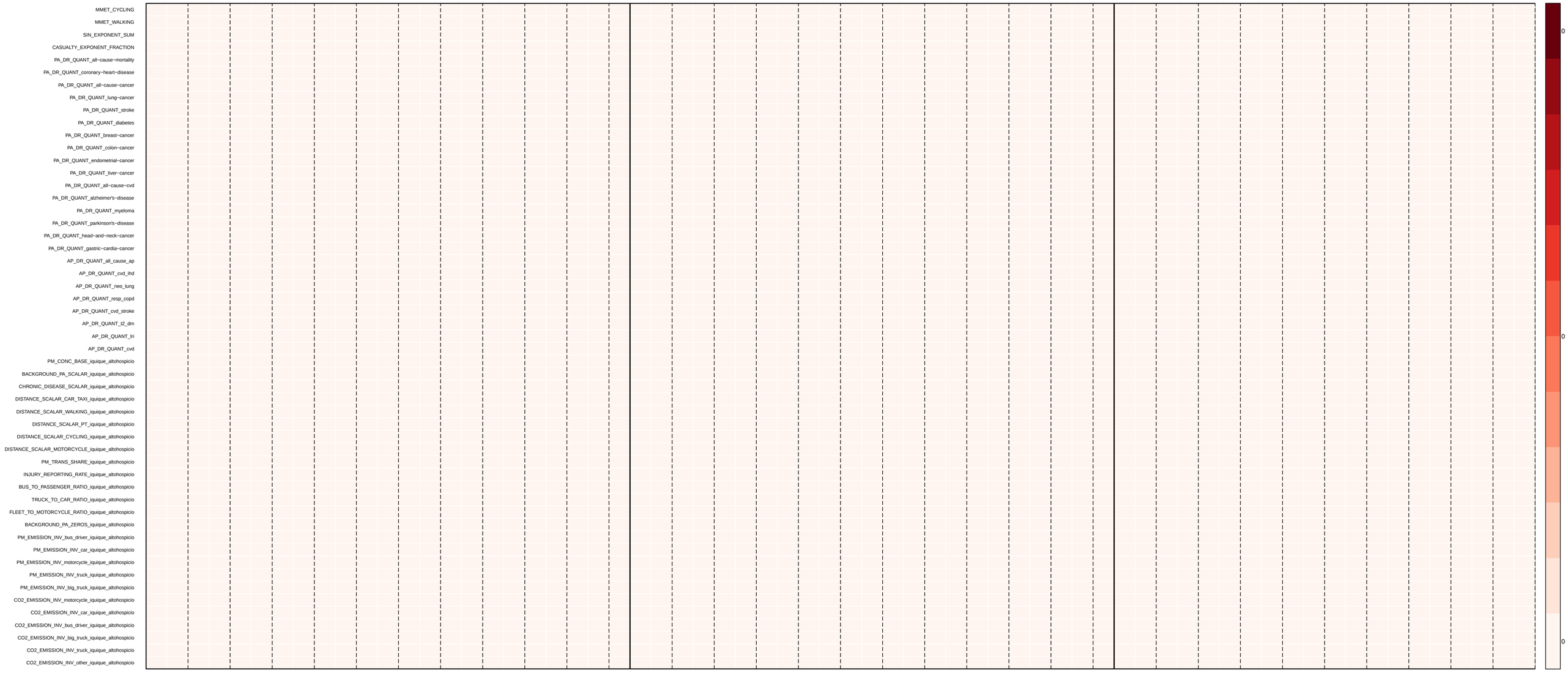
iquique_altohospicio − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



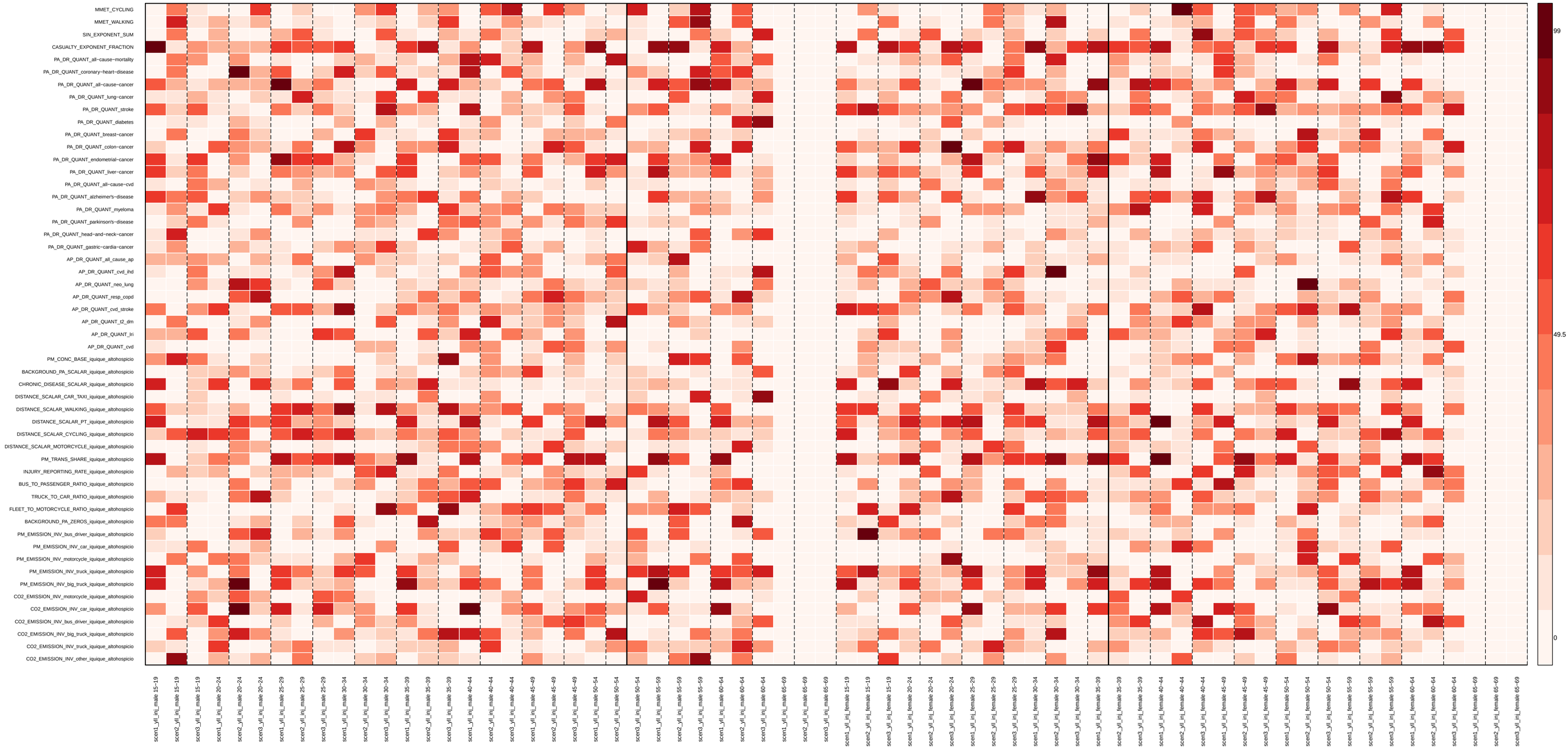
iquique_altohospicio − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



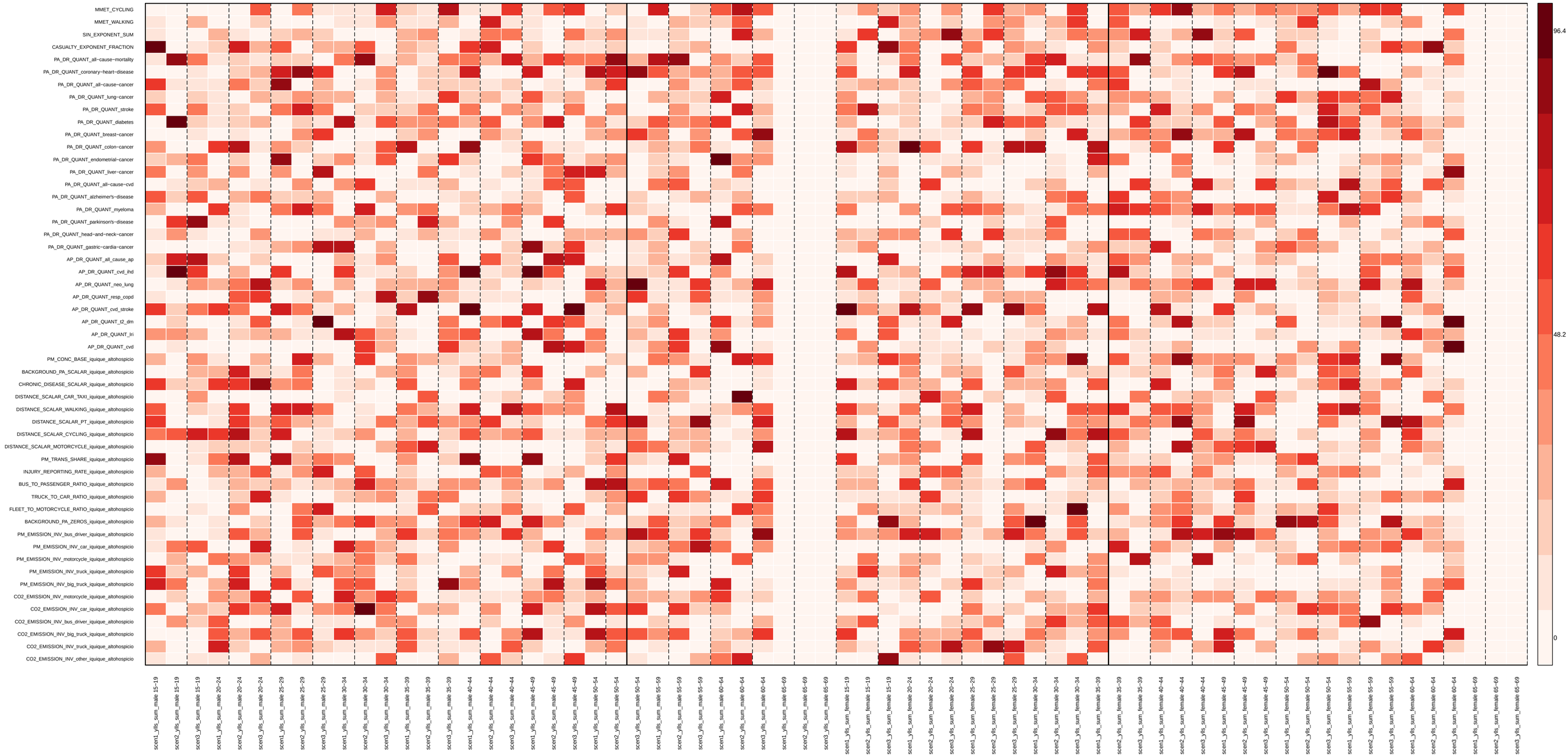
iquique_altohospicio − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



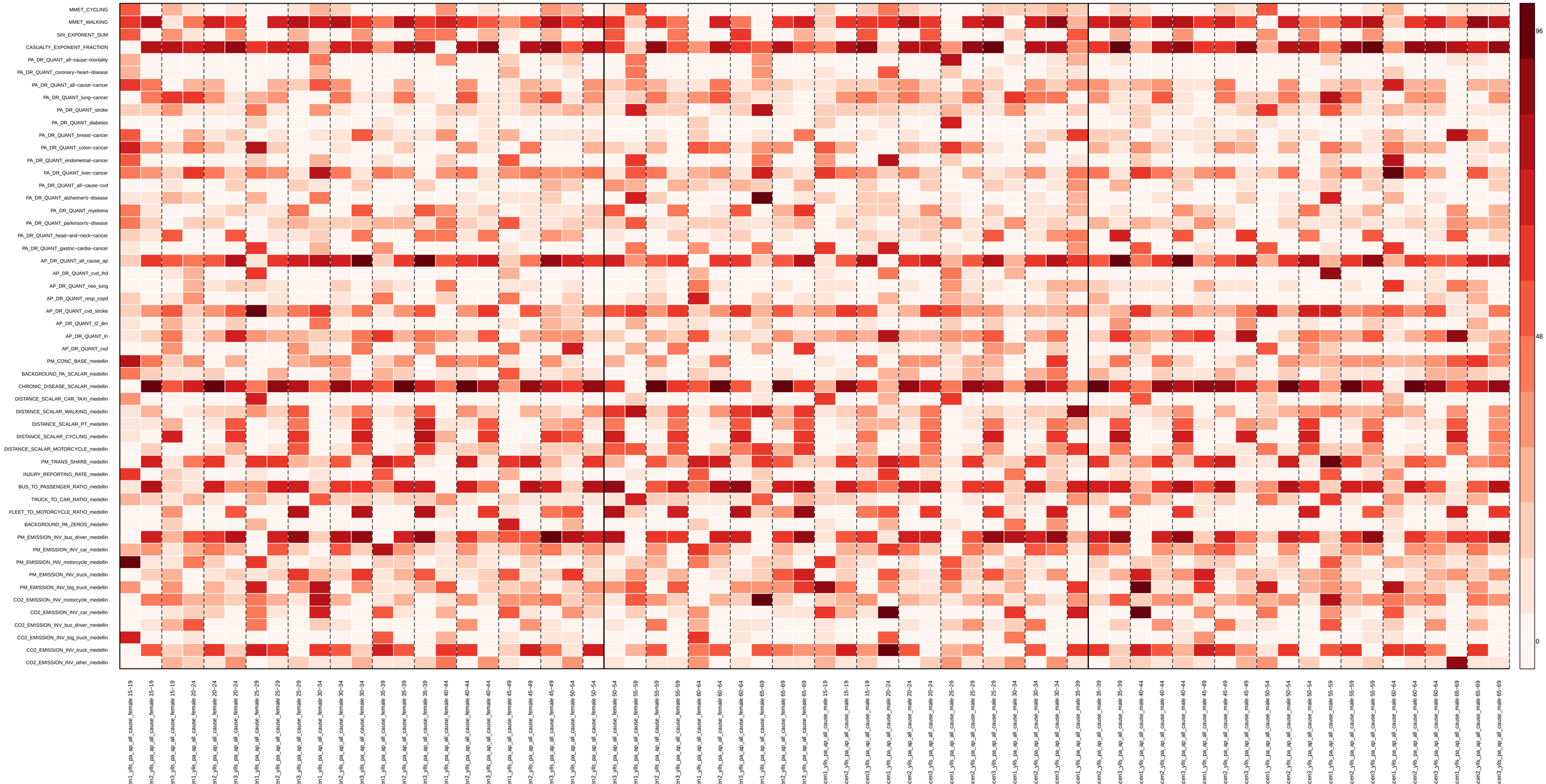
iquique_altohospicio − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



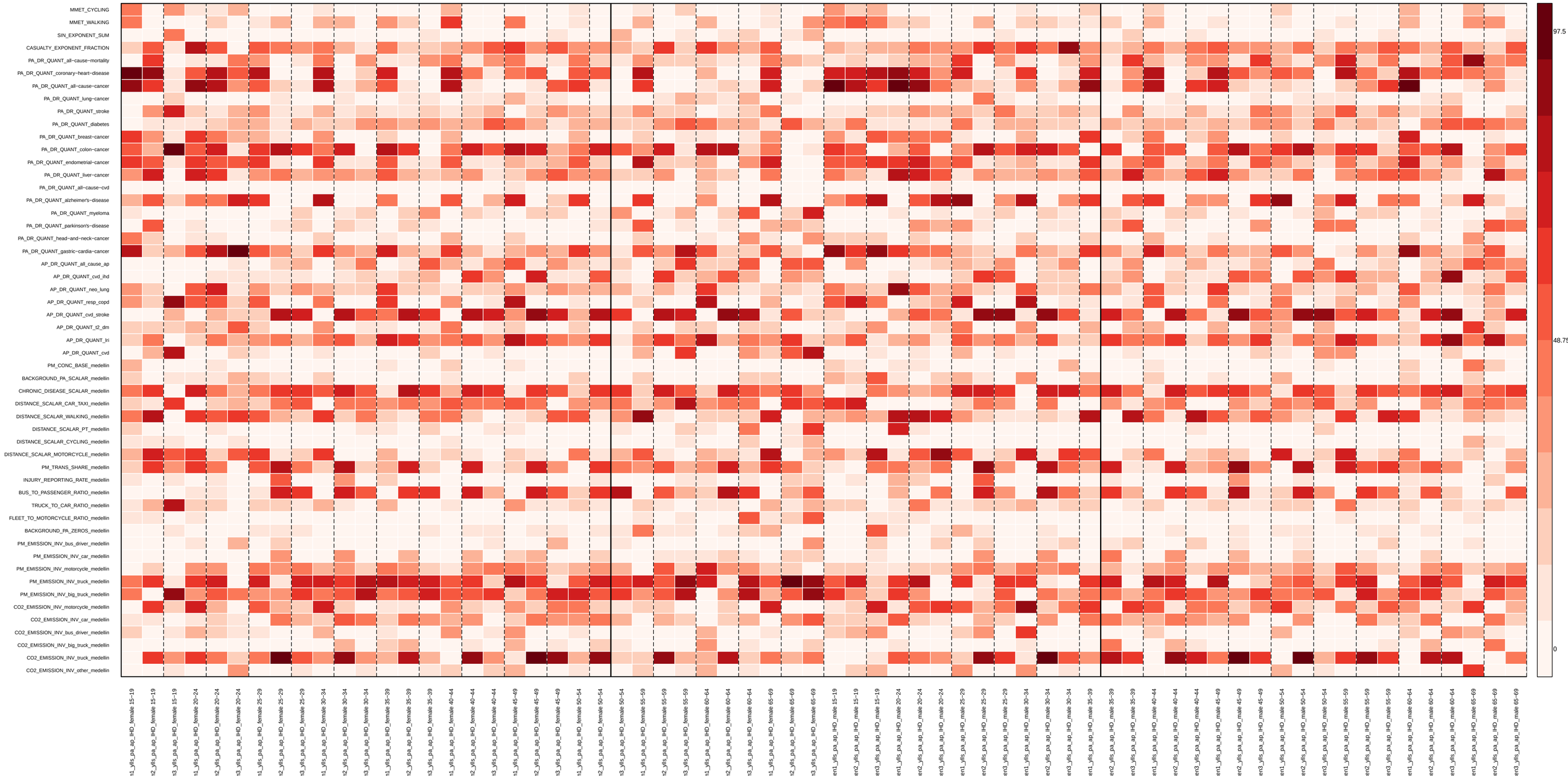
iquique_altohospicio − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



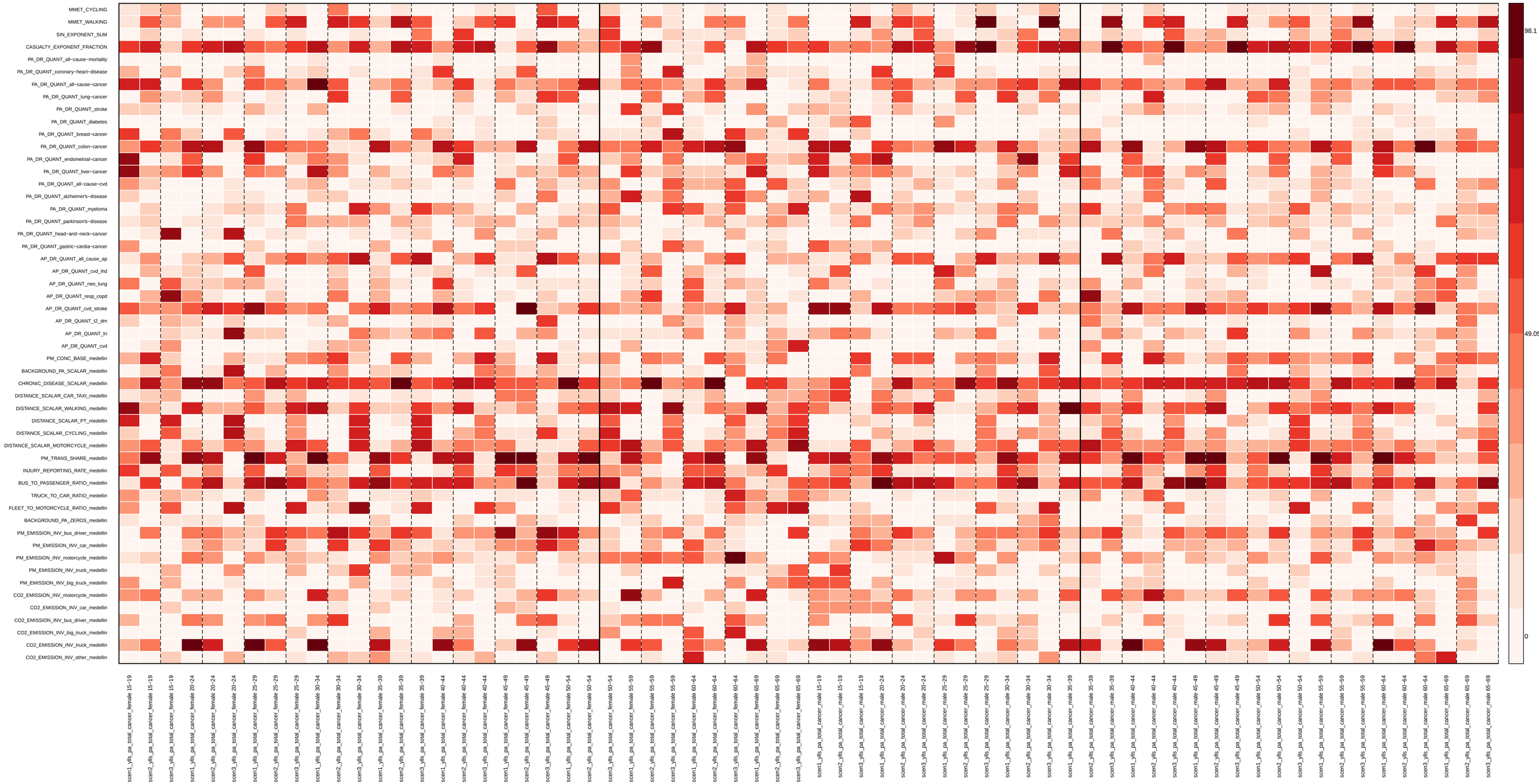
medellin − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



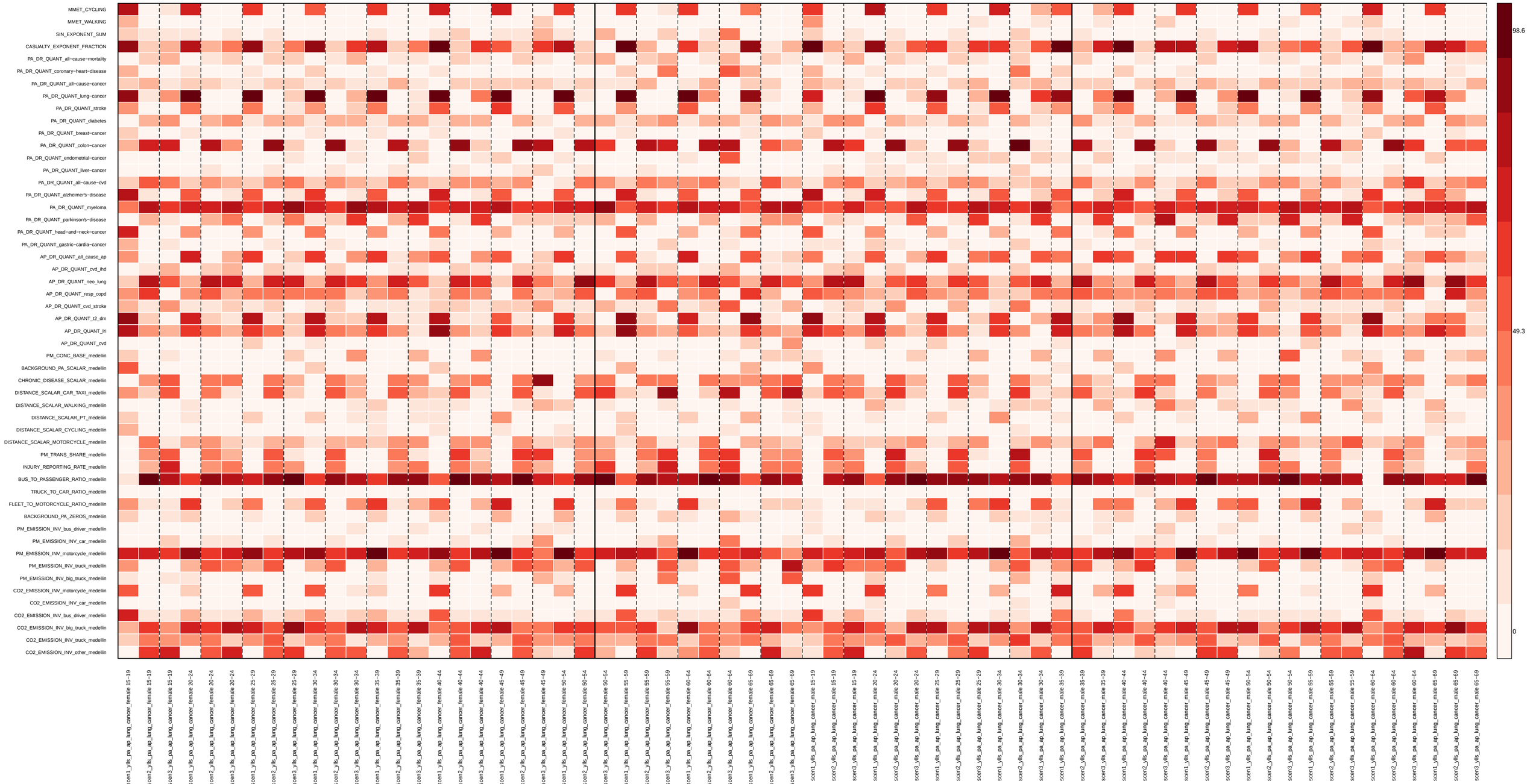
medellin − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



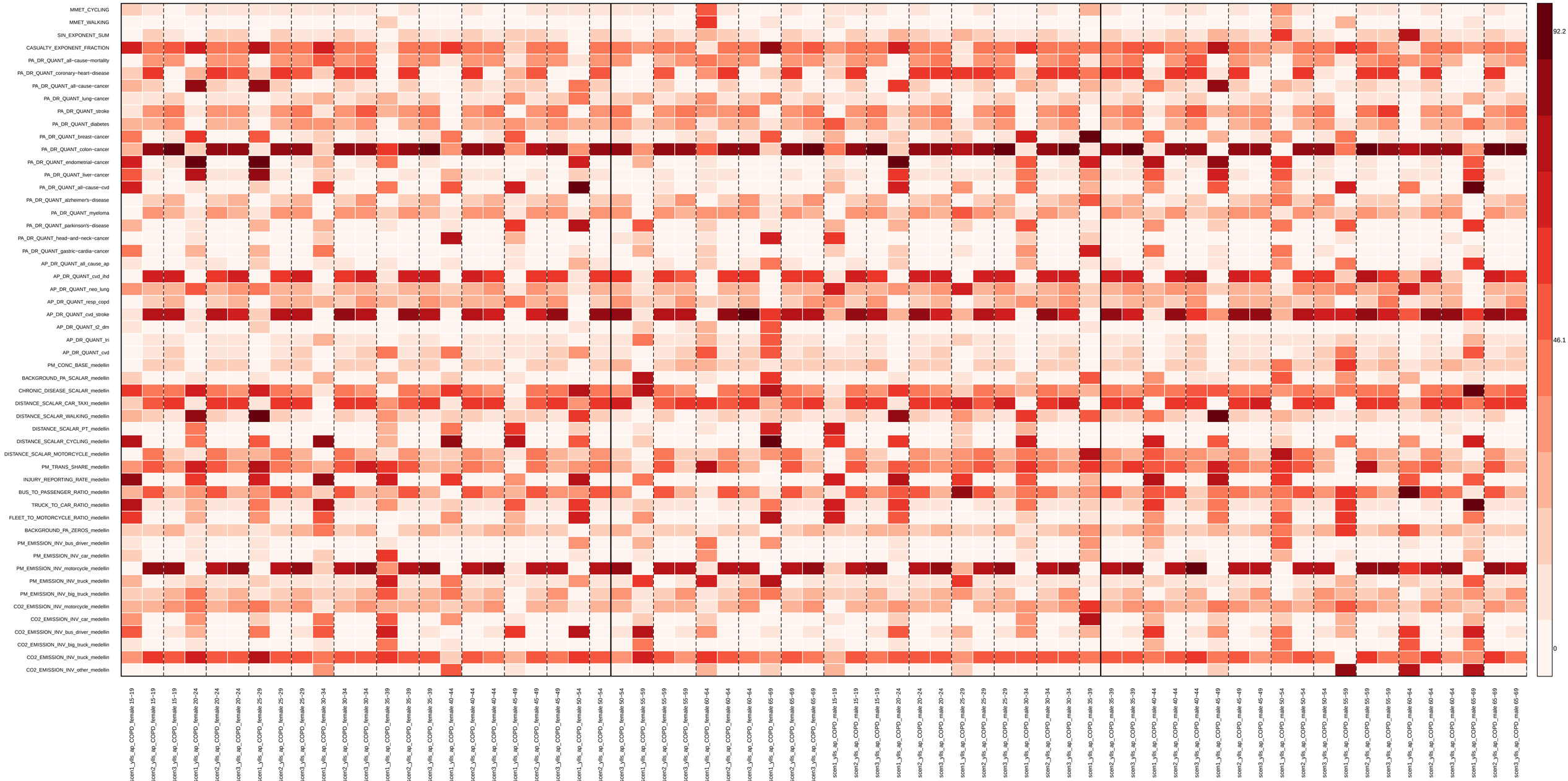
medellin − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



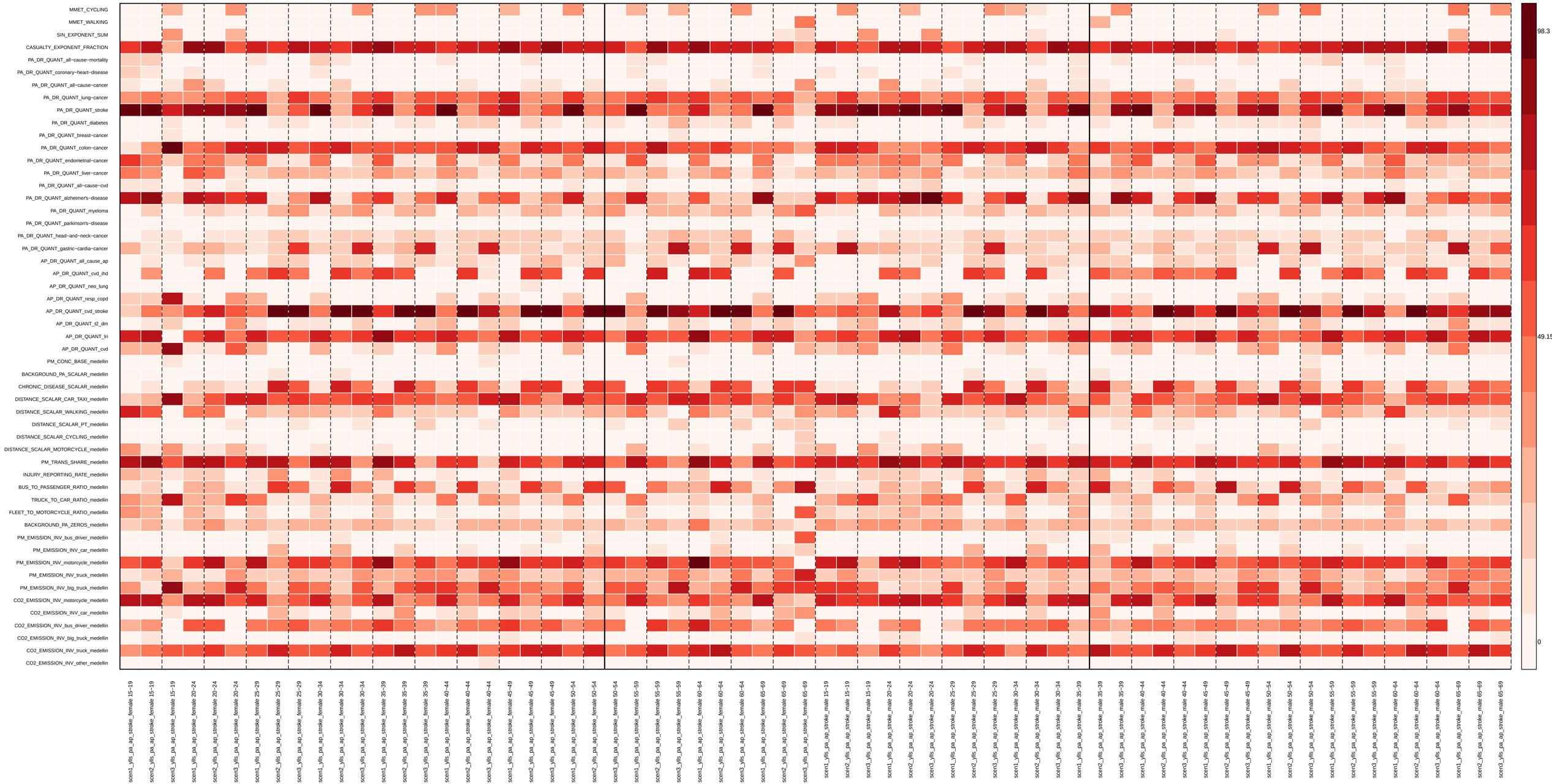
medellin − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



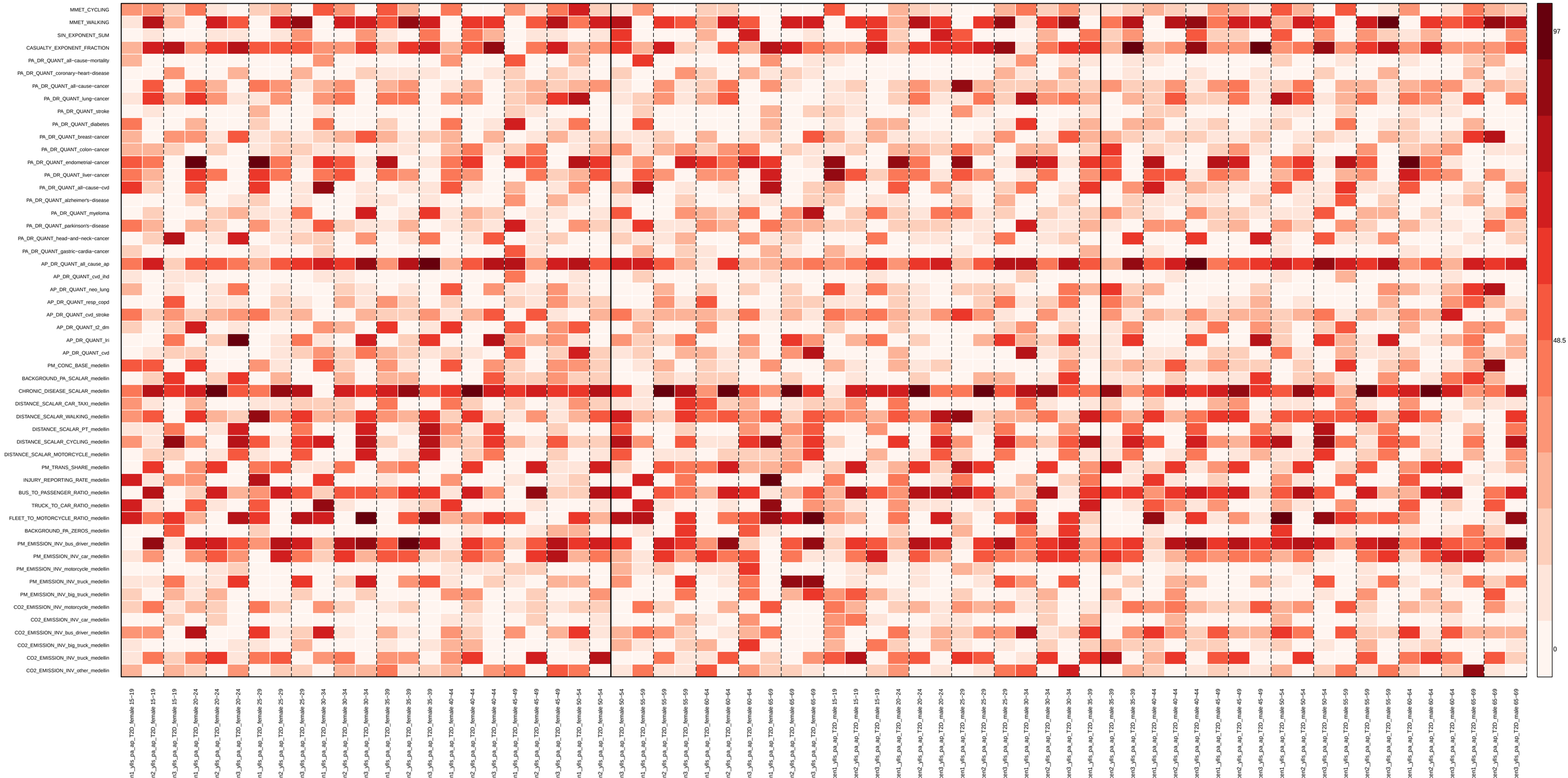
medellin − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



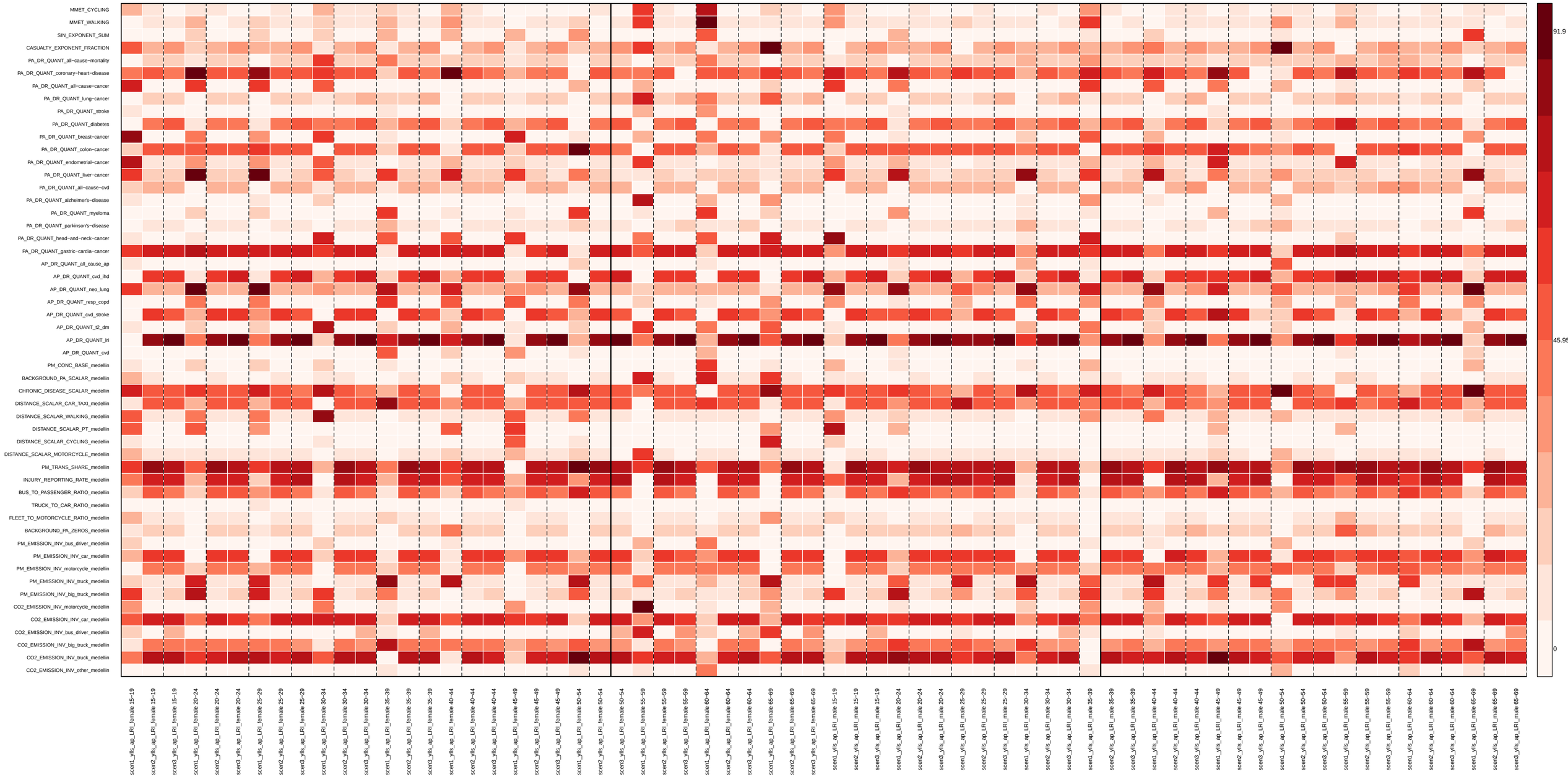
medellin − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



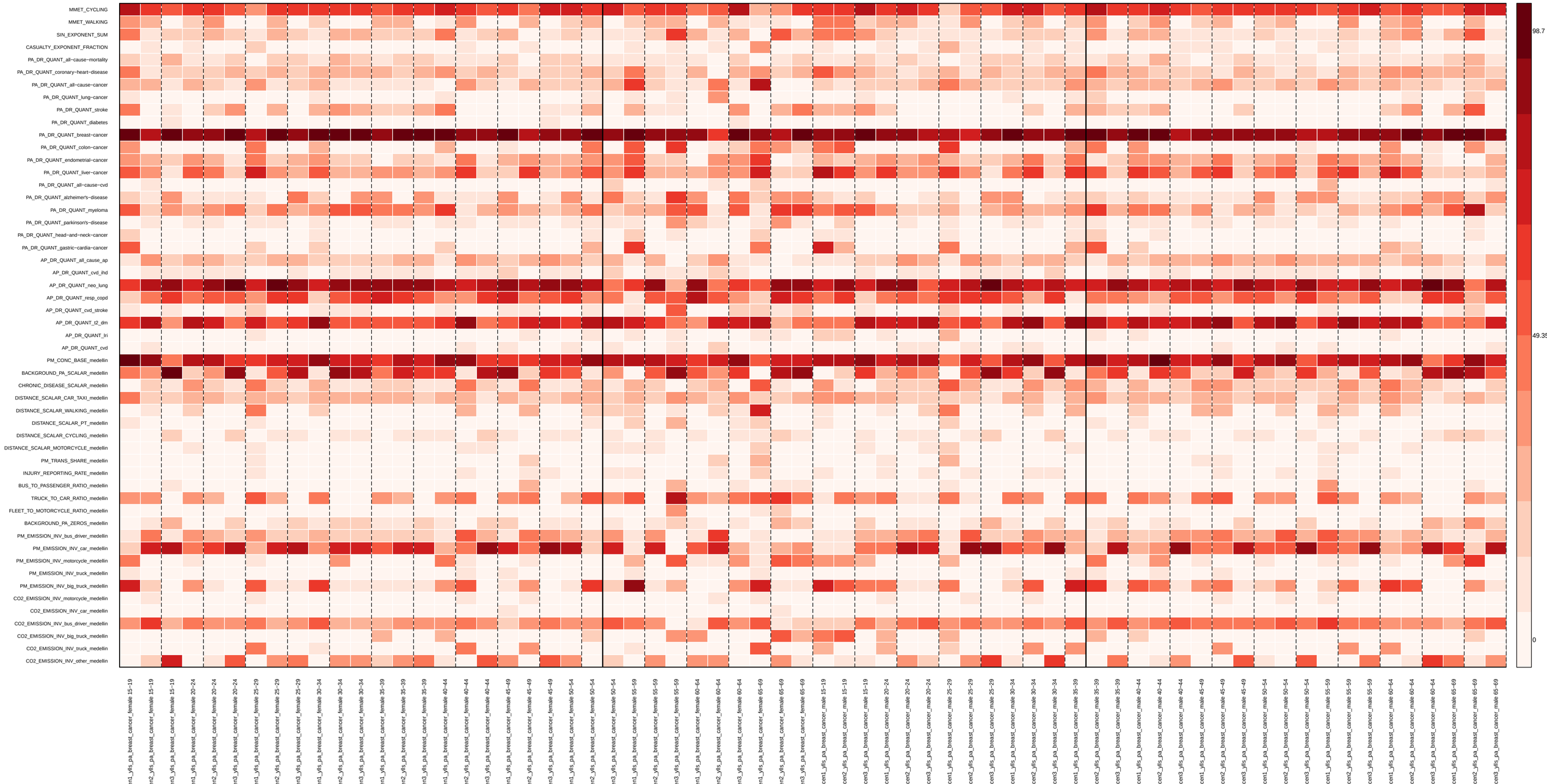
medellin − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



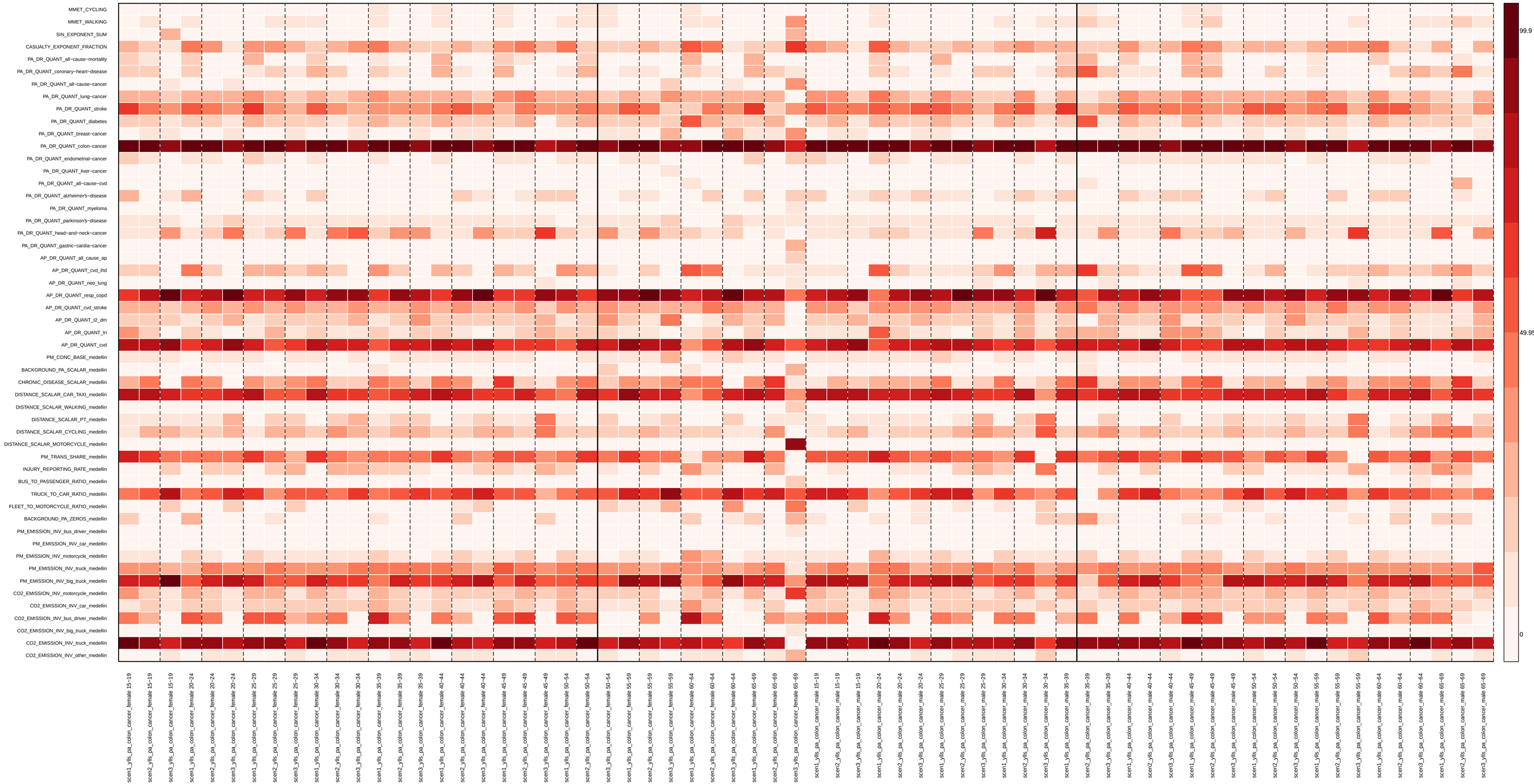
medellin − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



medellin − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

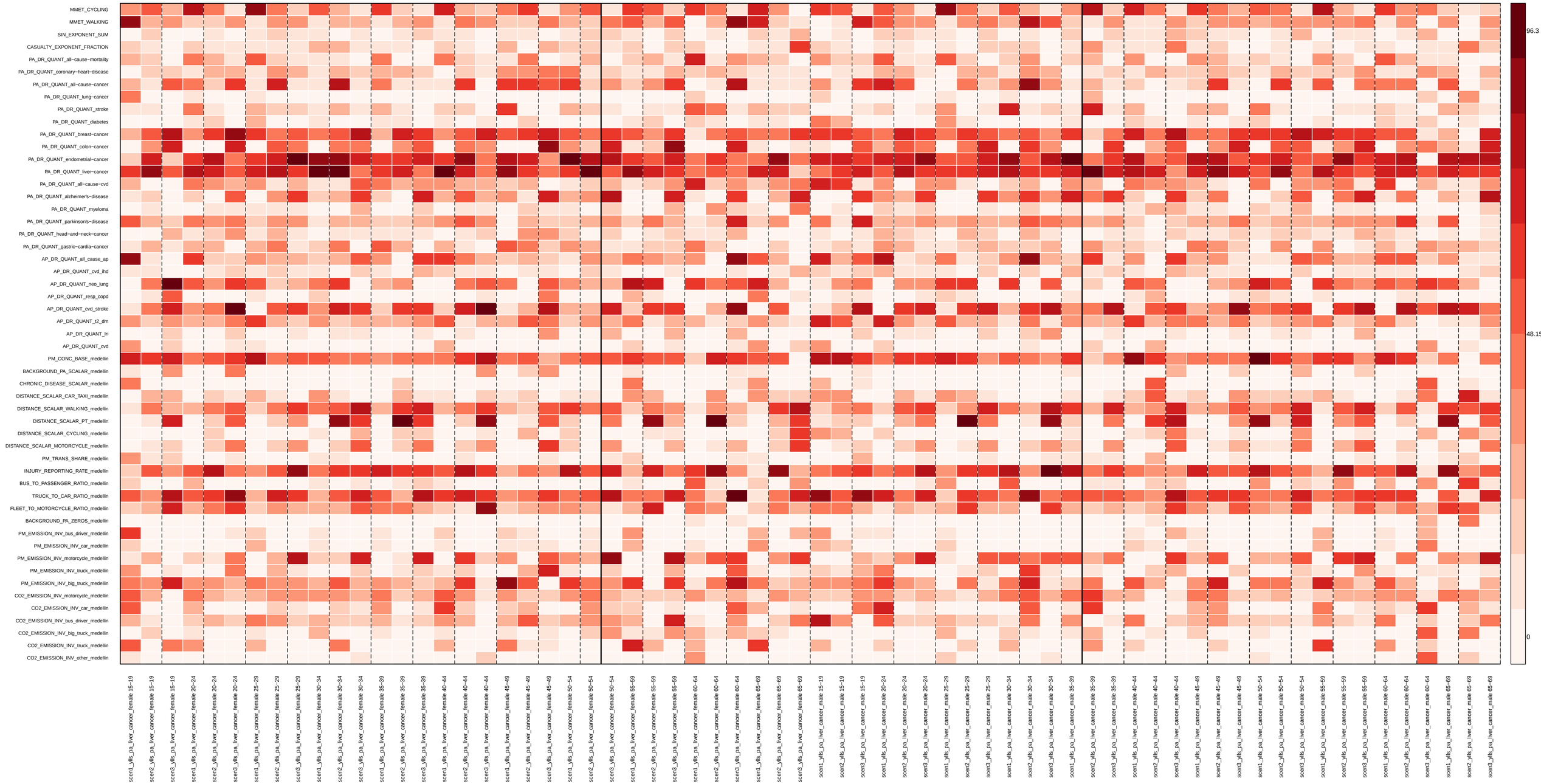


medellin − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

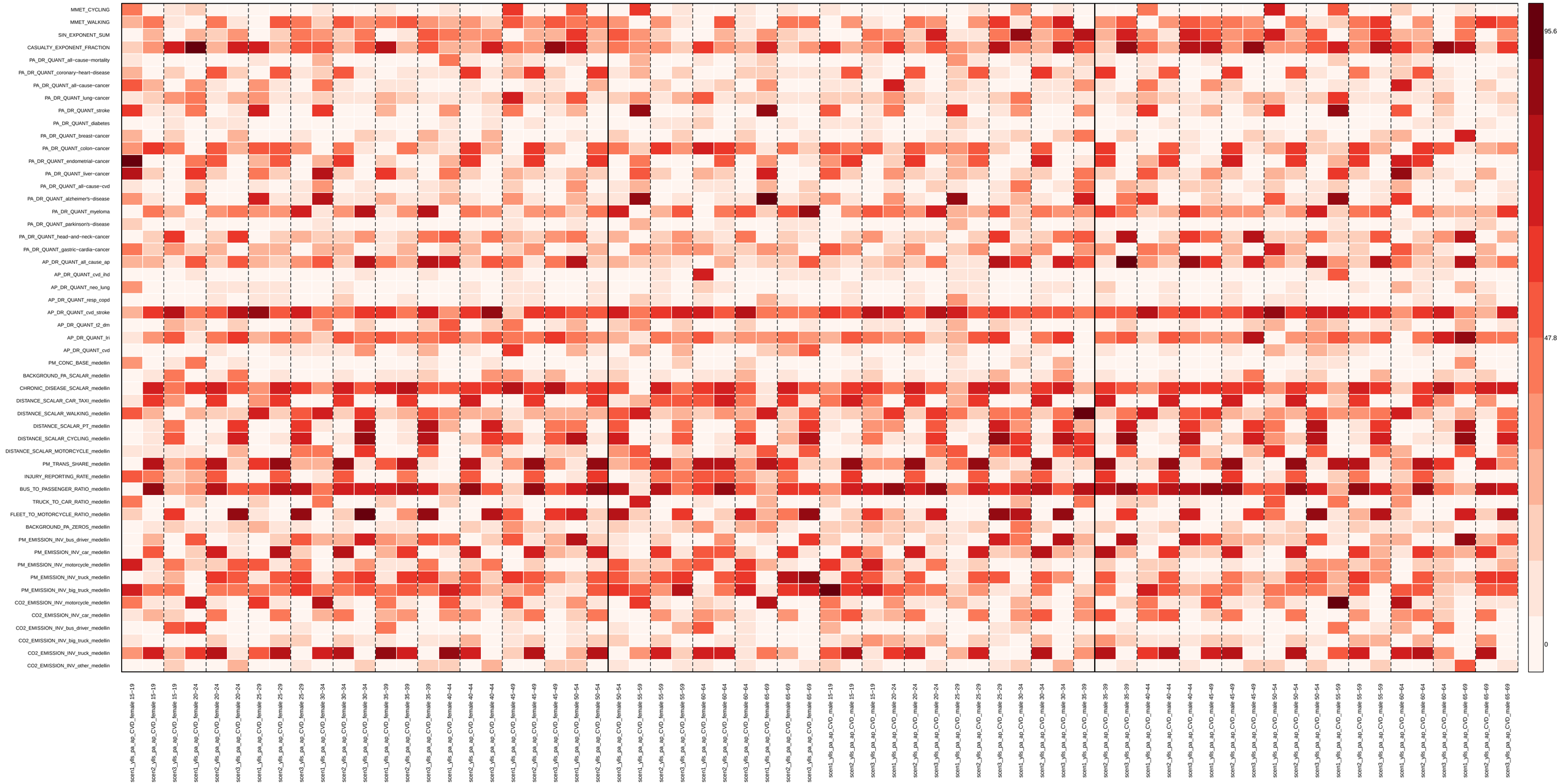


medellin − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

medellin − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

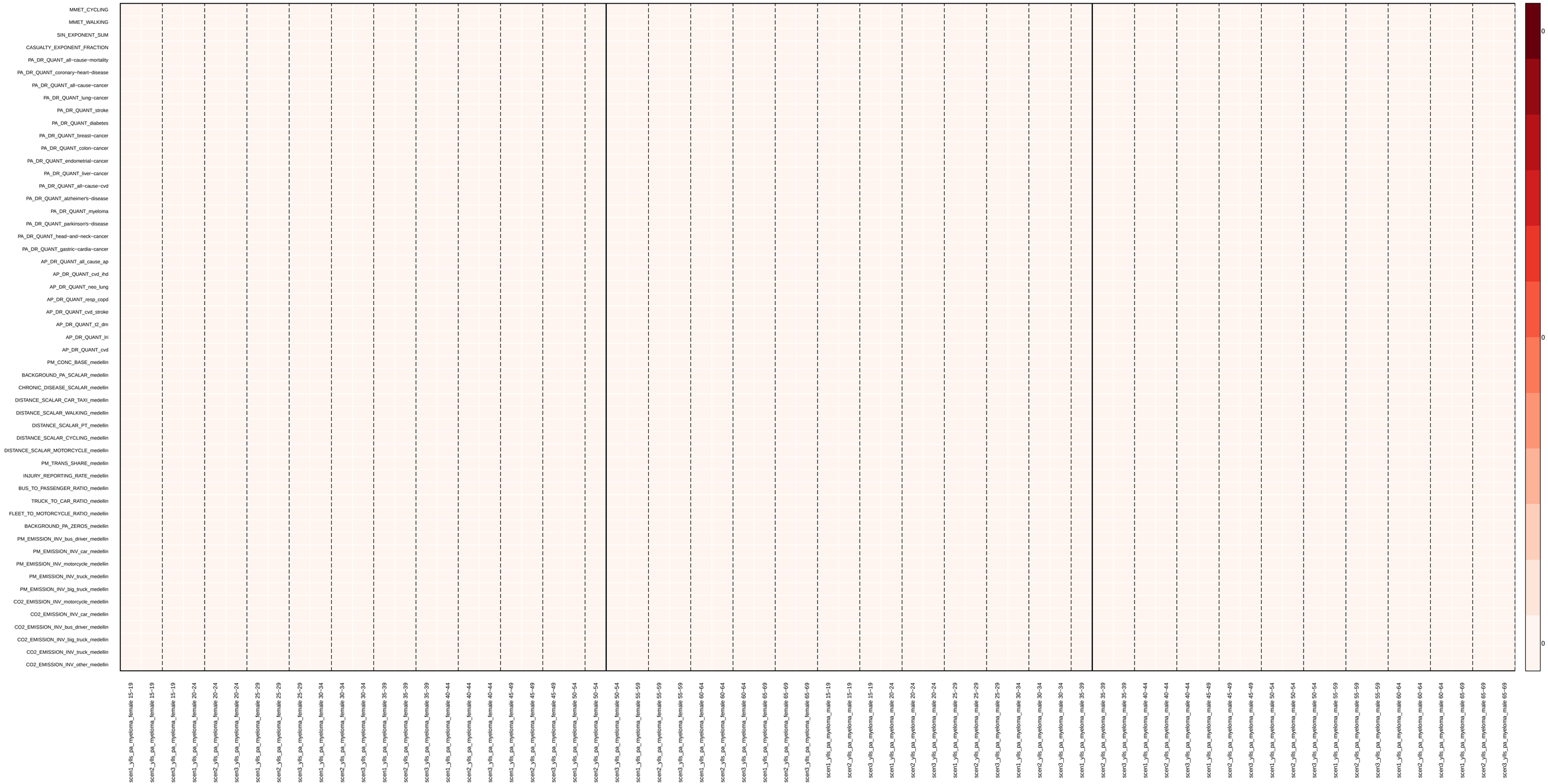


medellin − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

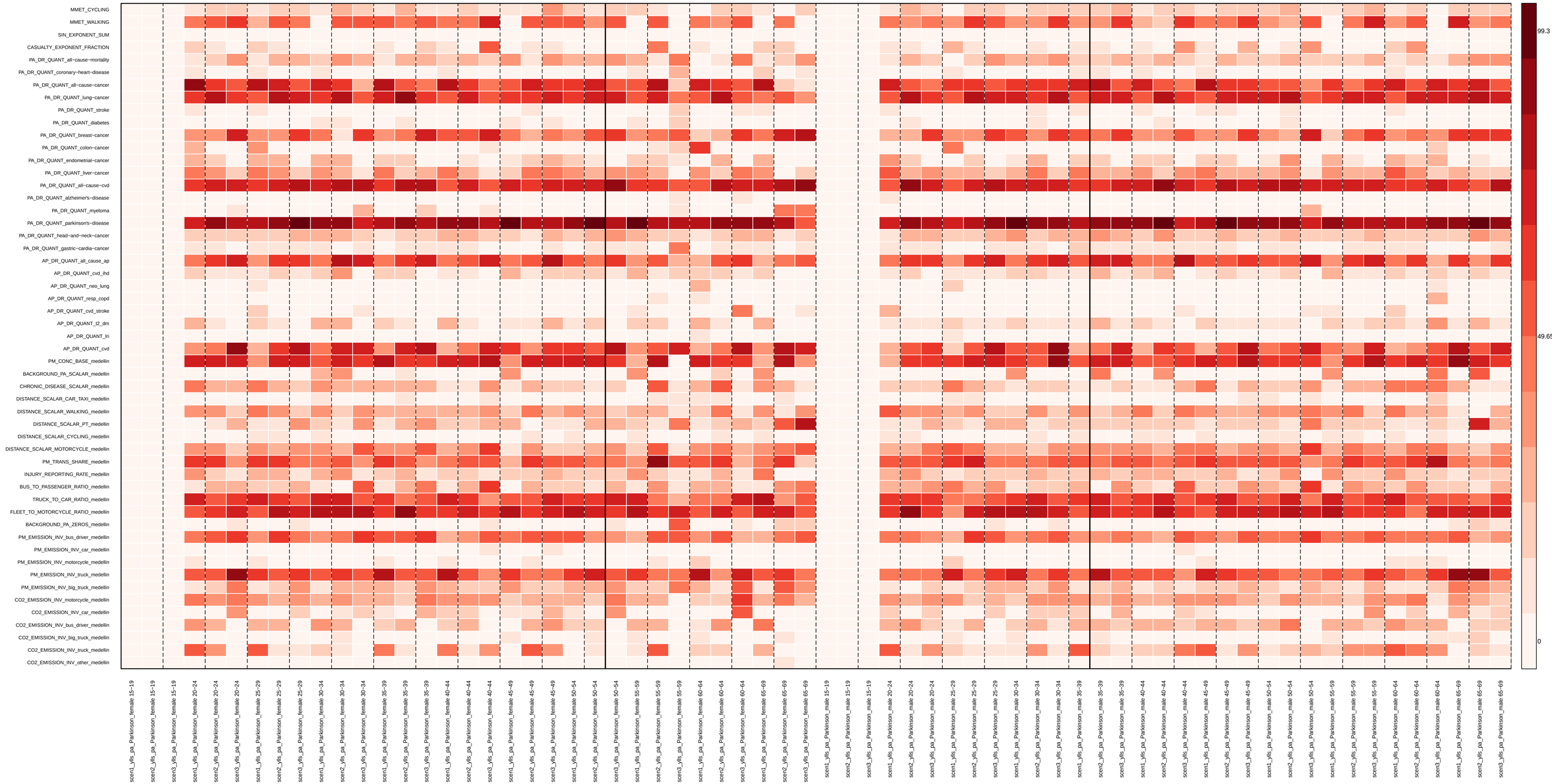


medellin − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

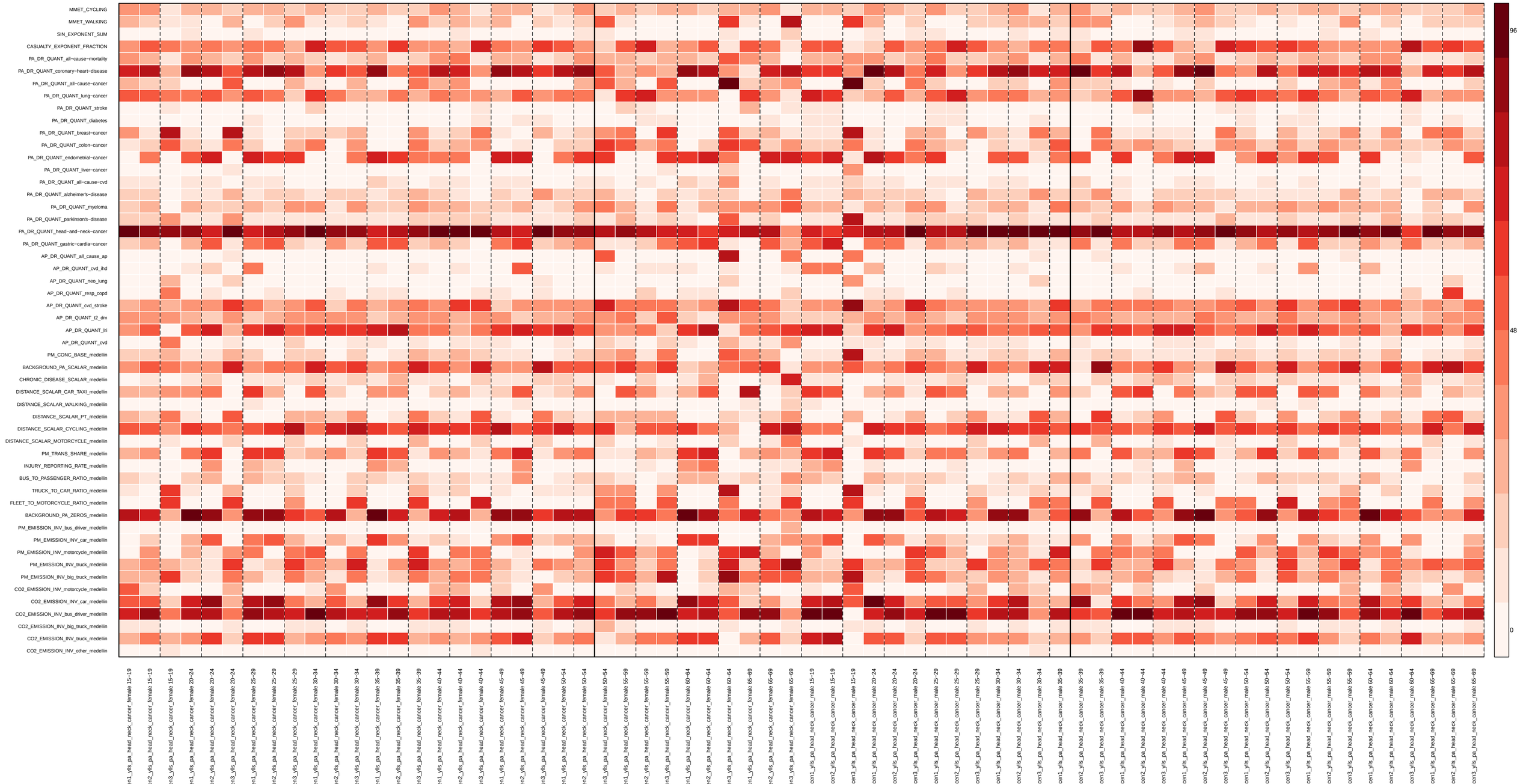
medellin − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



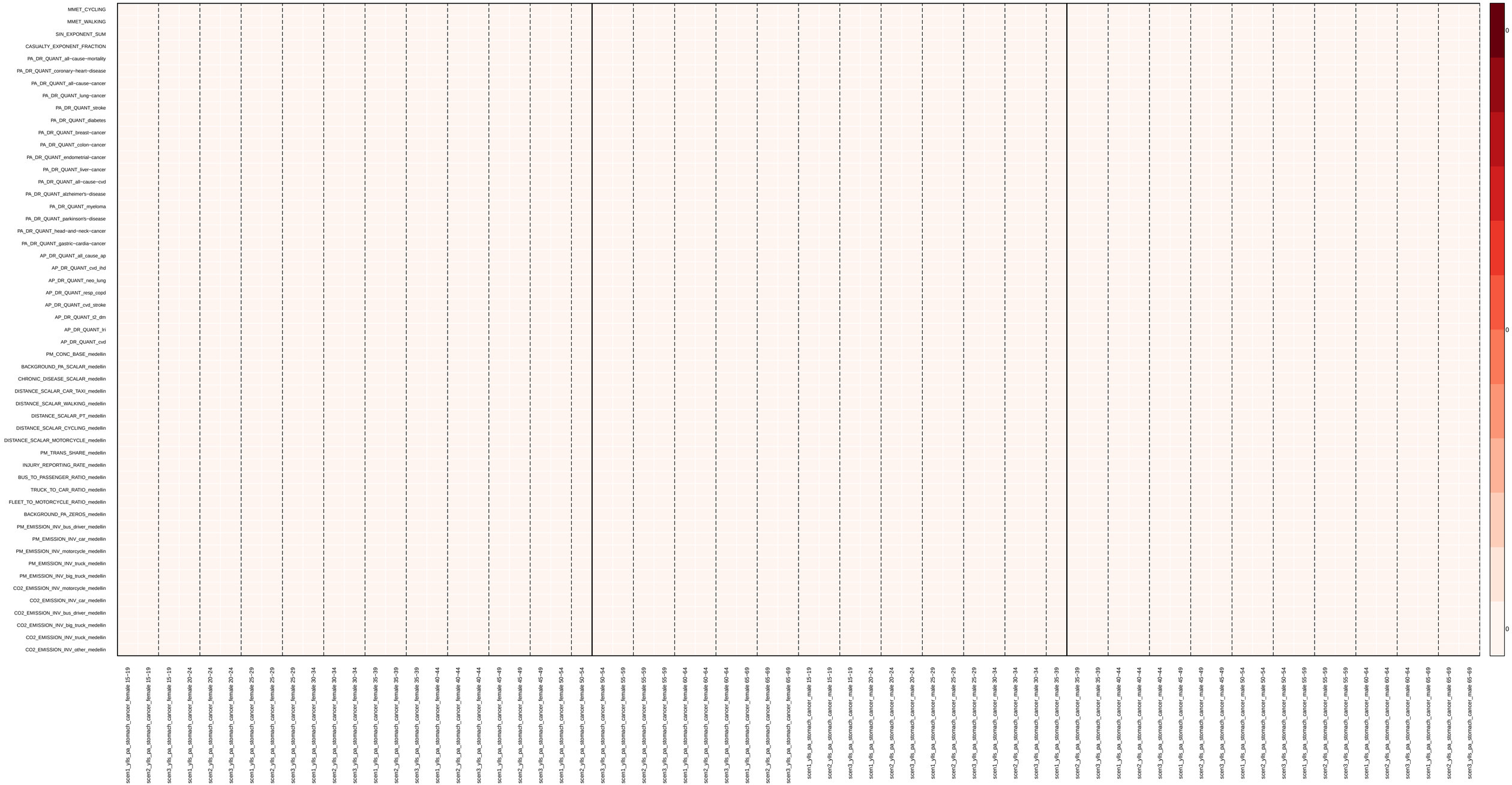
medellin − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



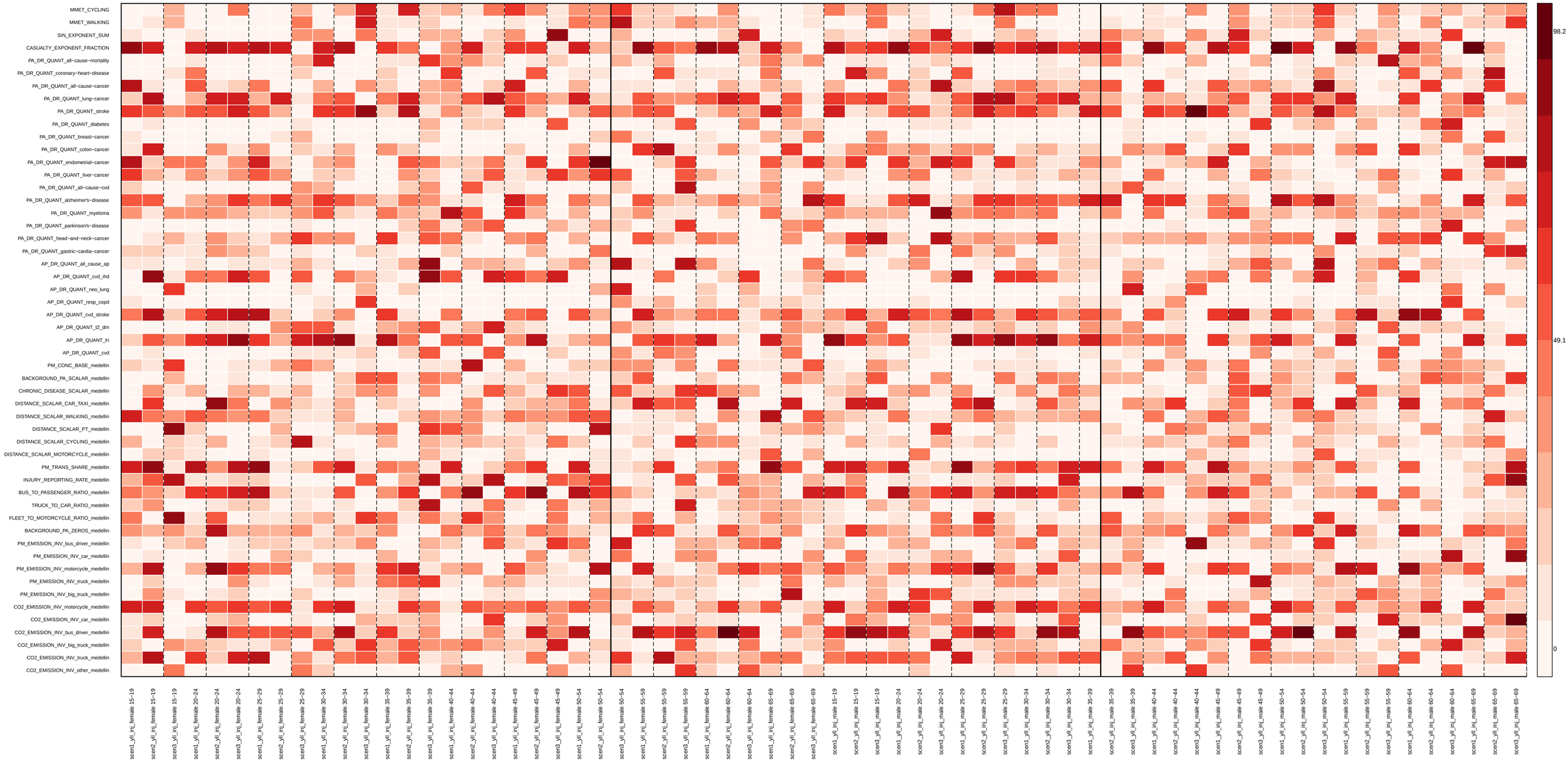
medellin − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



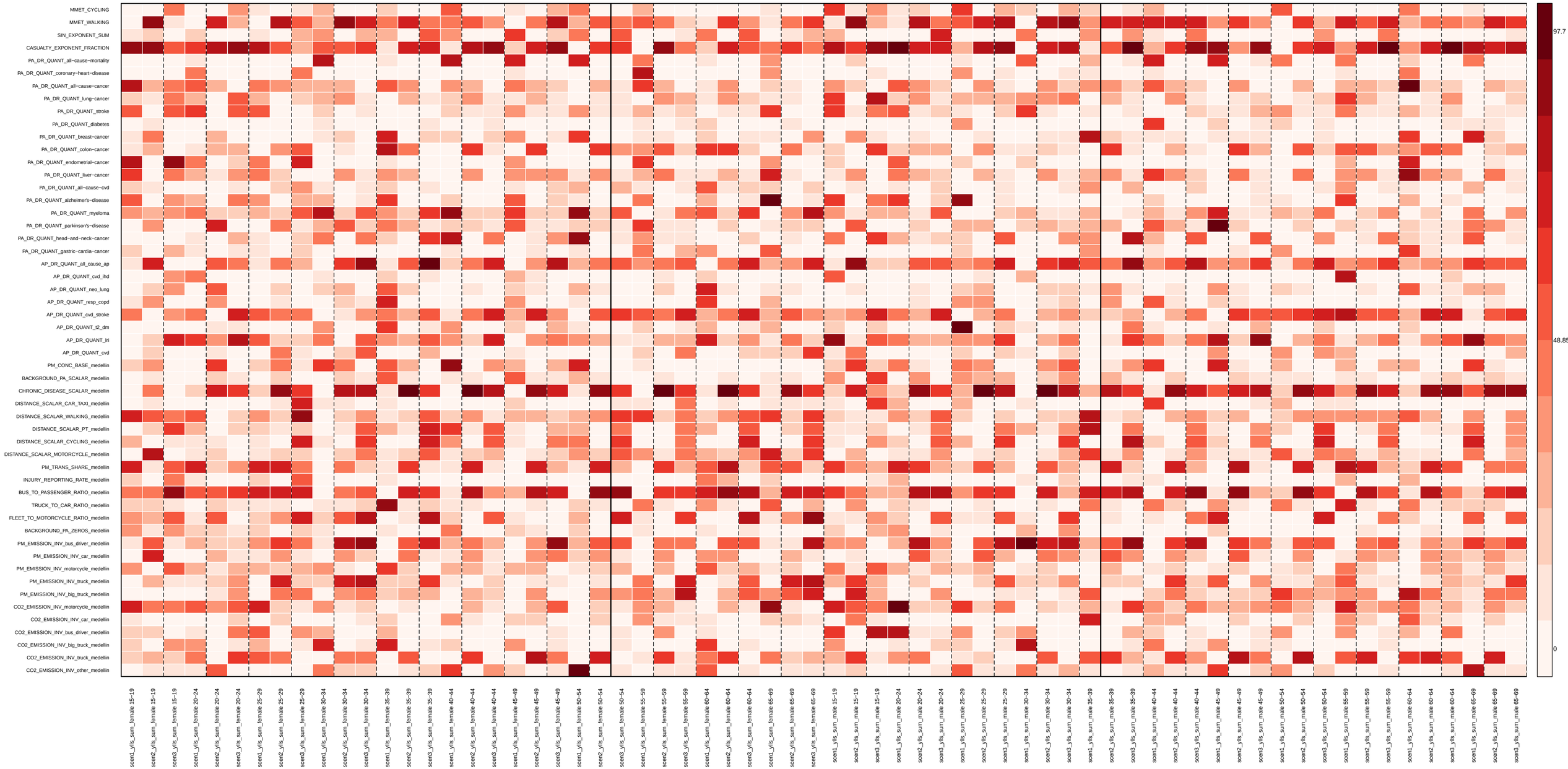
medellin − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



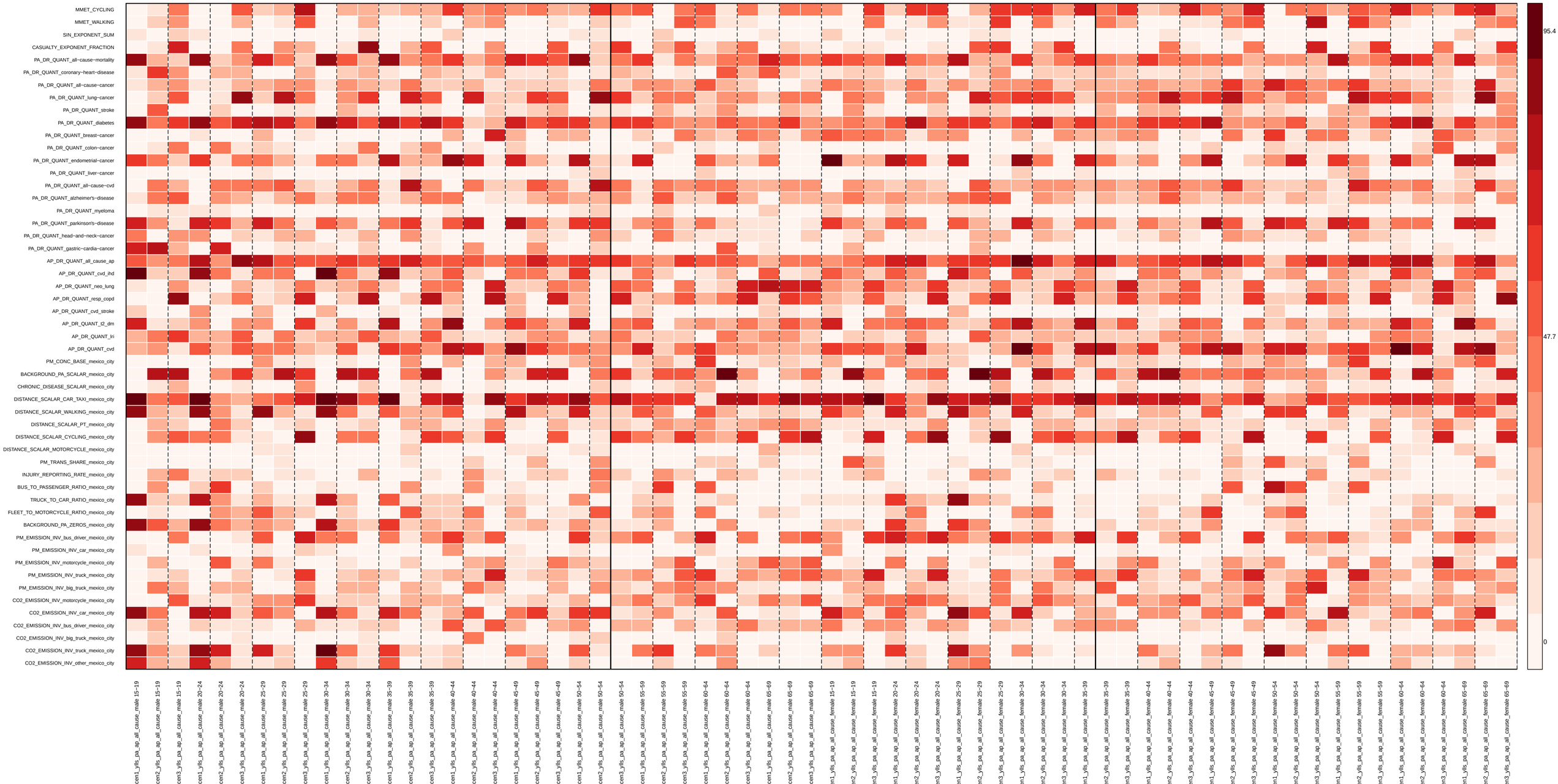
medellin − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



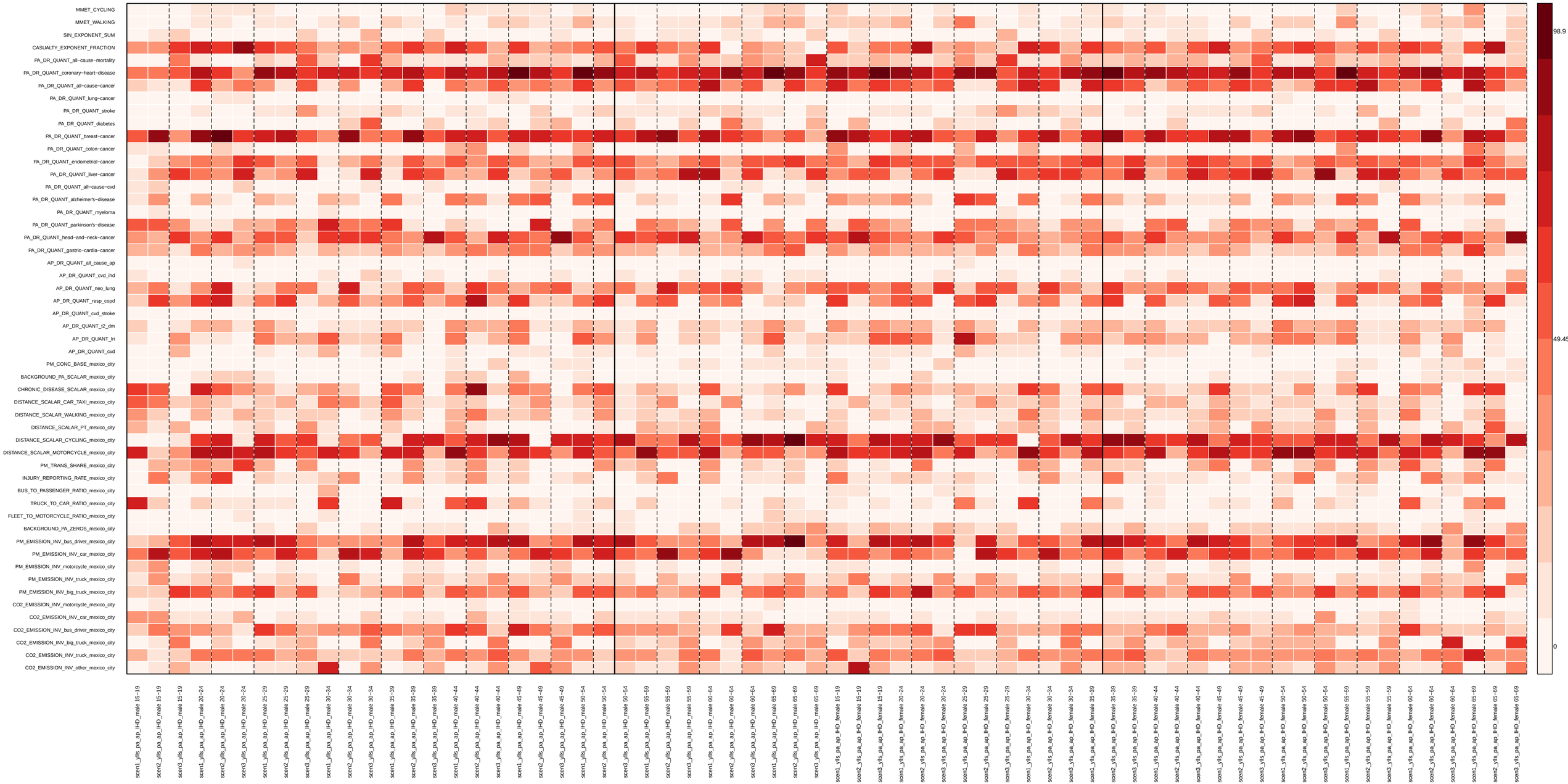
medellin − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



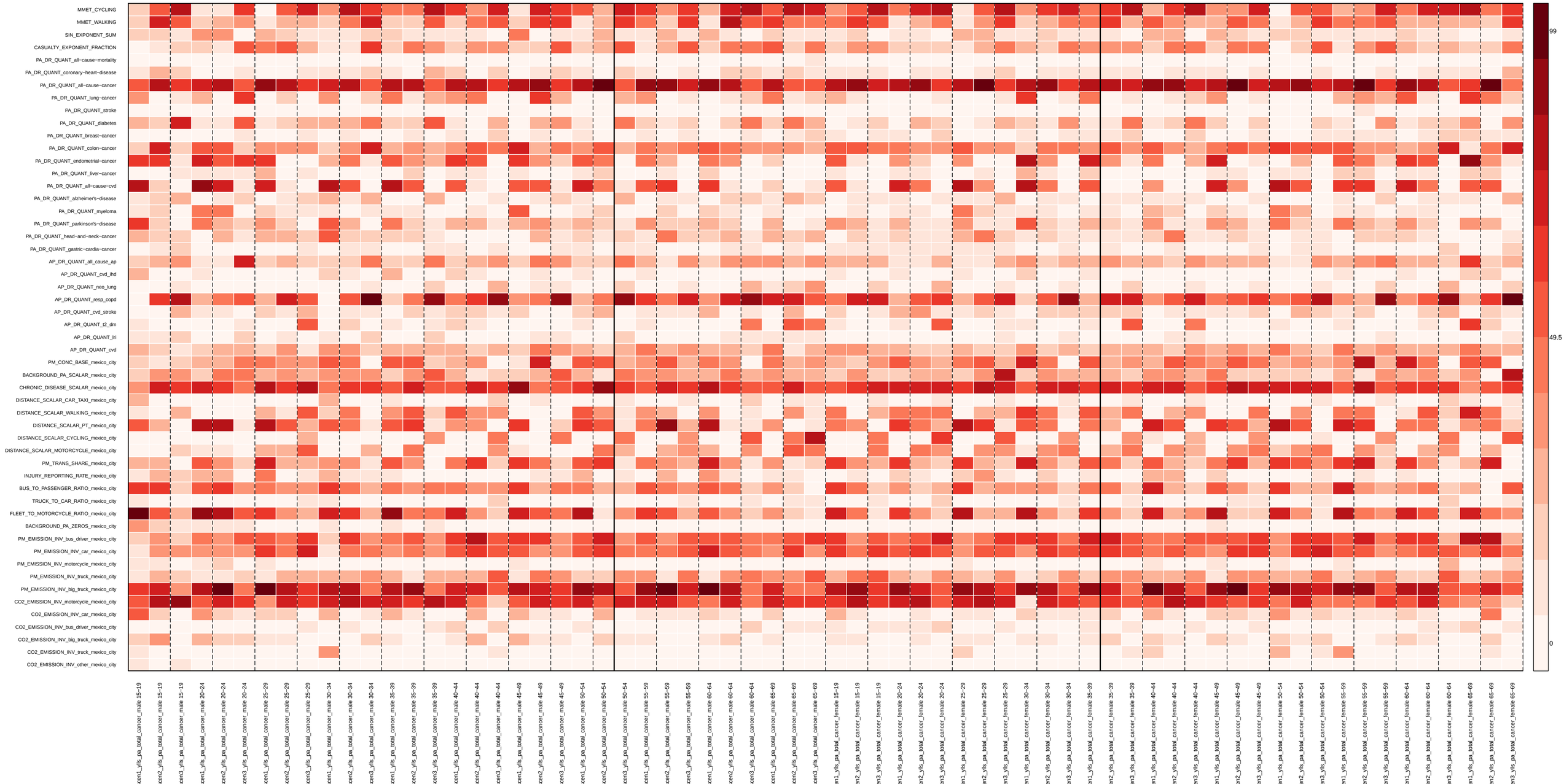
mexico_city − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



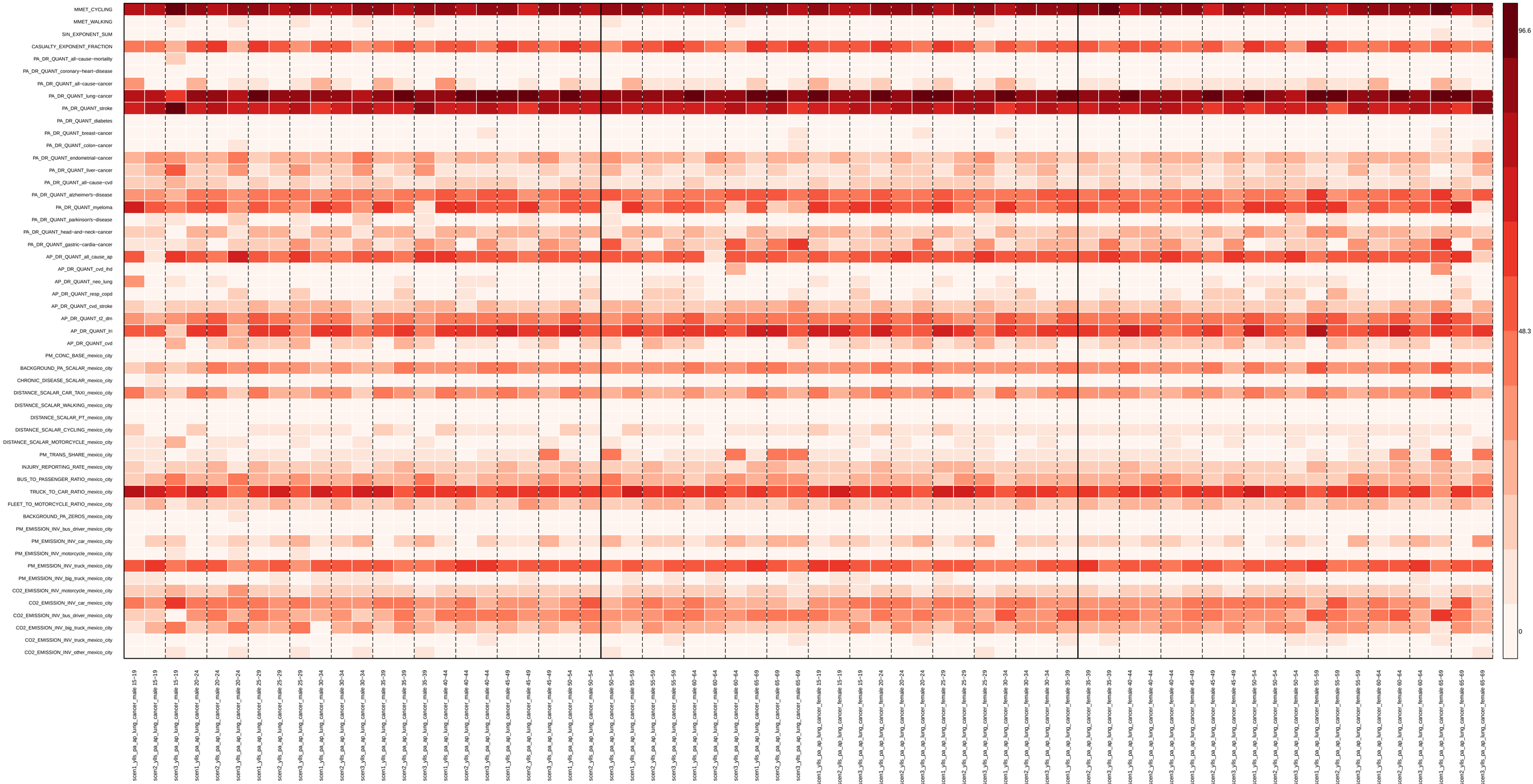
mexico_city − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



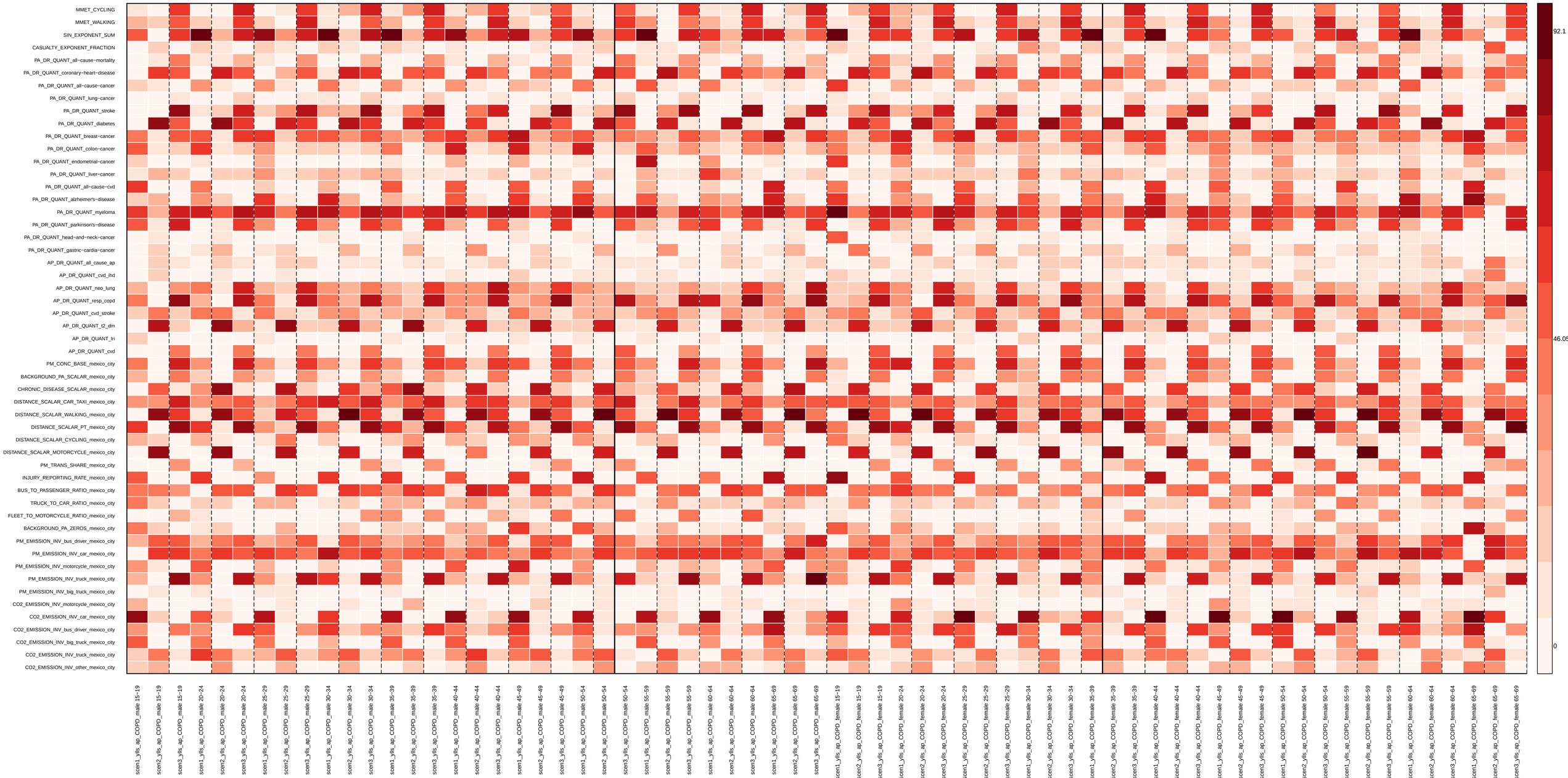
mexico_city − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



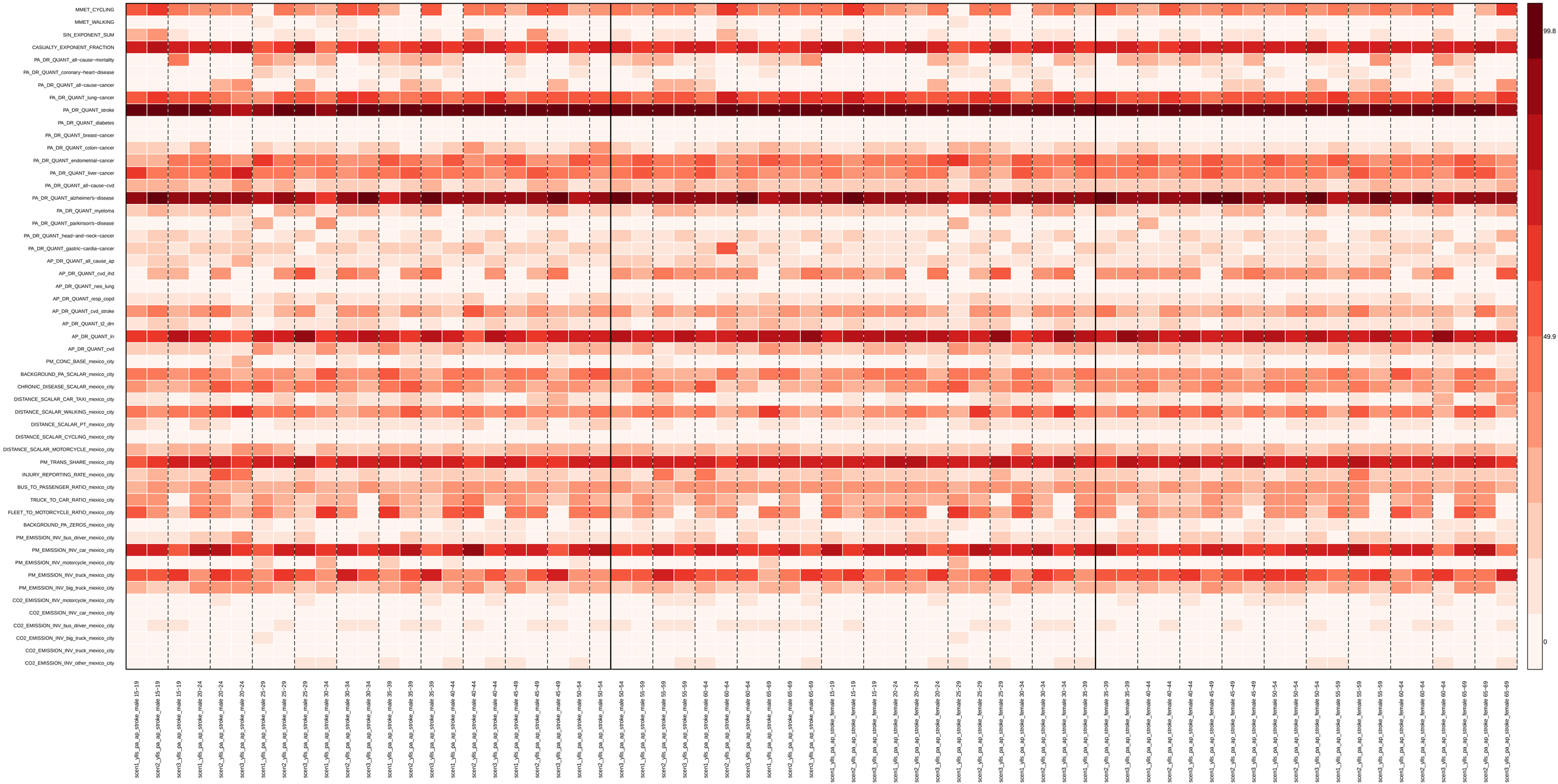
mexico_city − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



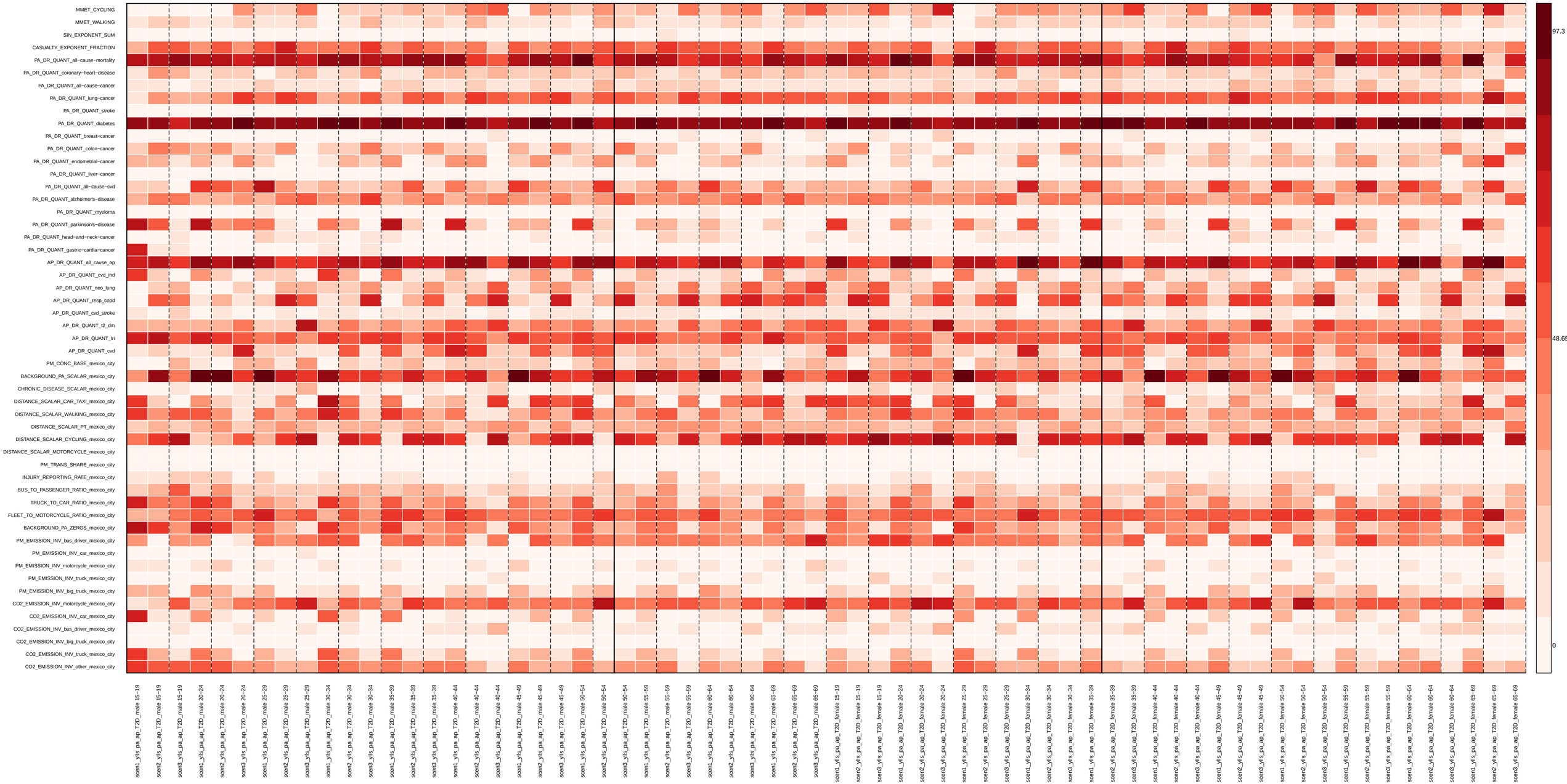
mexico_city − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



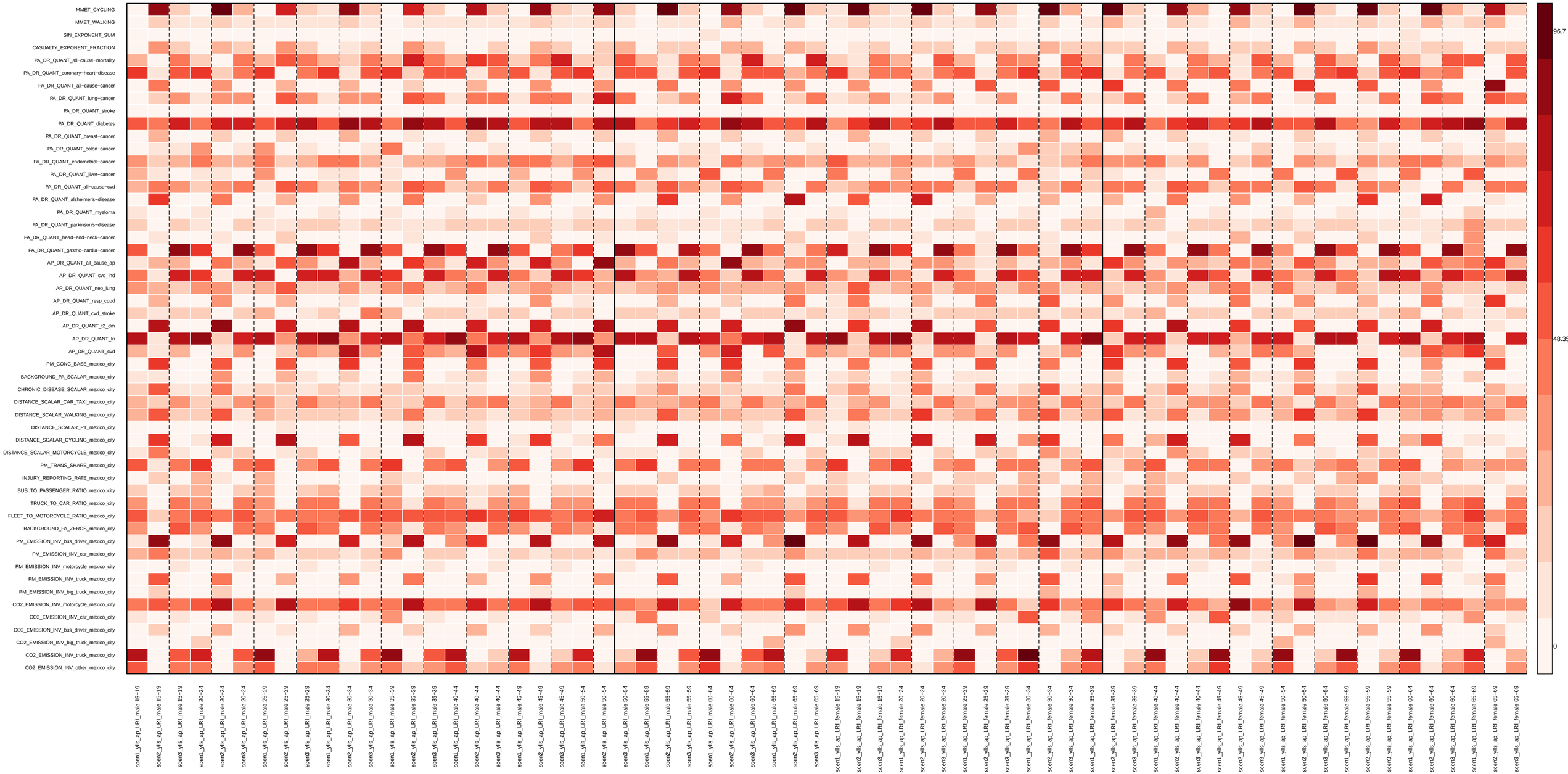
mexico_city − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



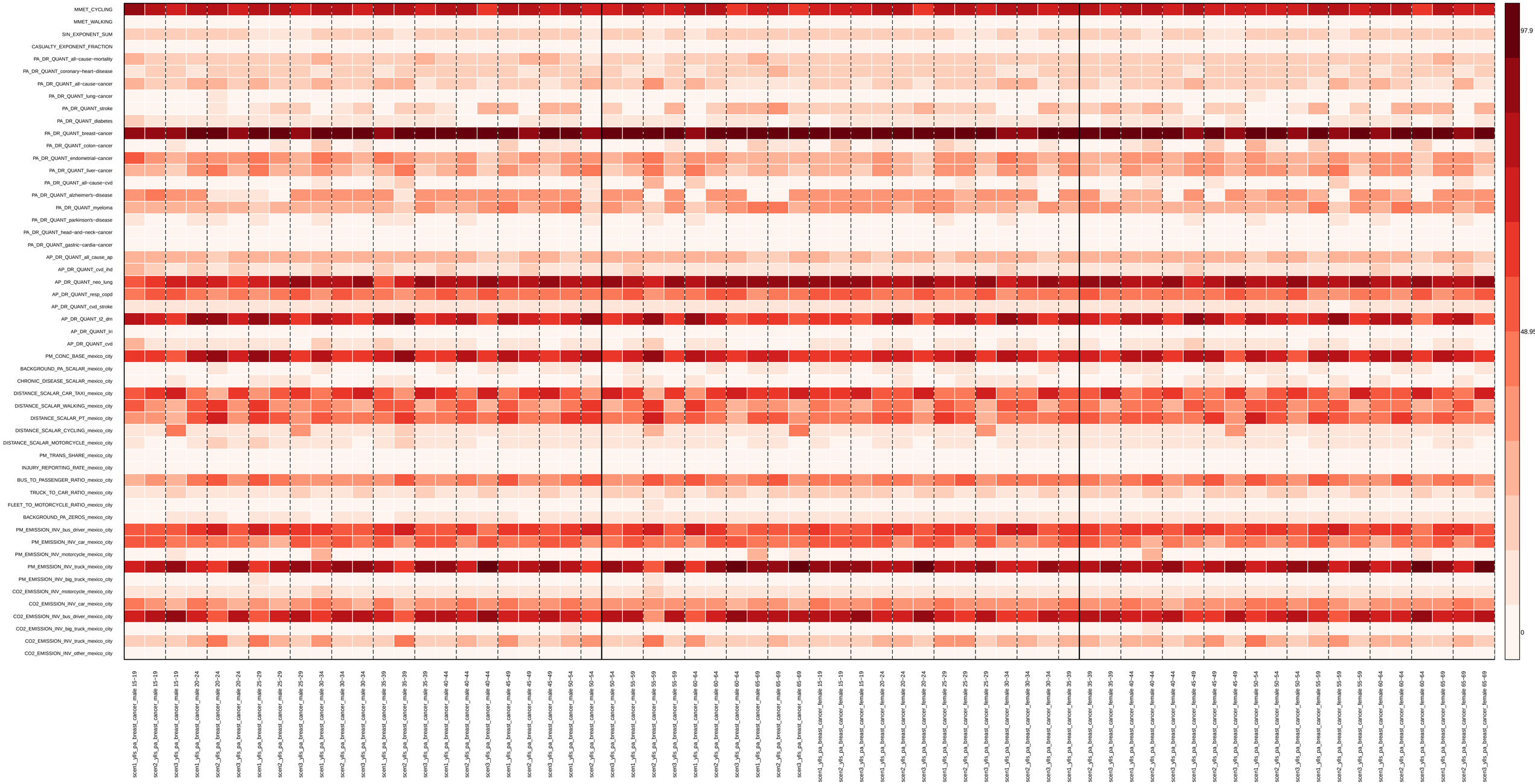
mexico_city − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



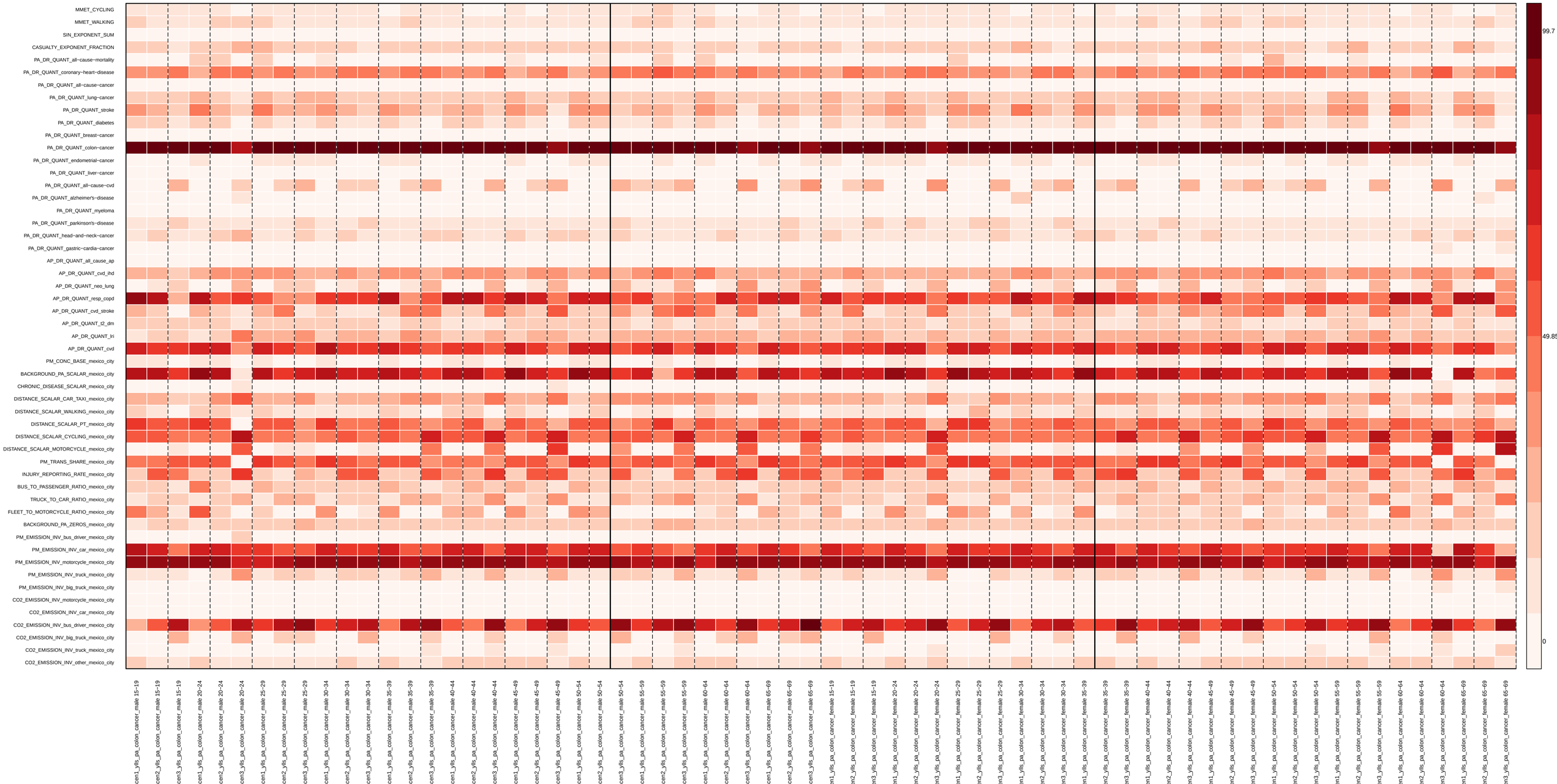
mexico_city − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



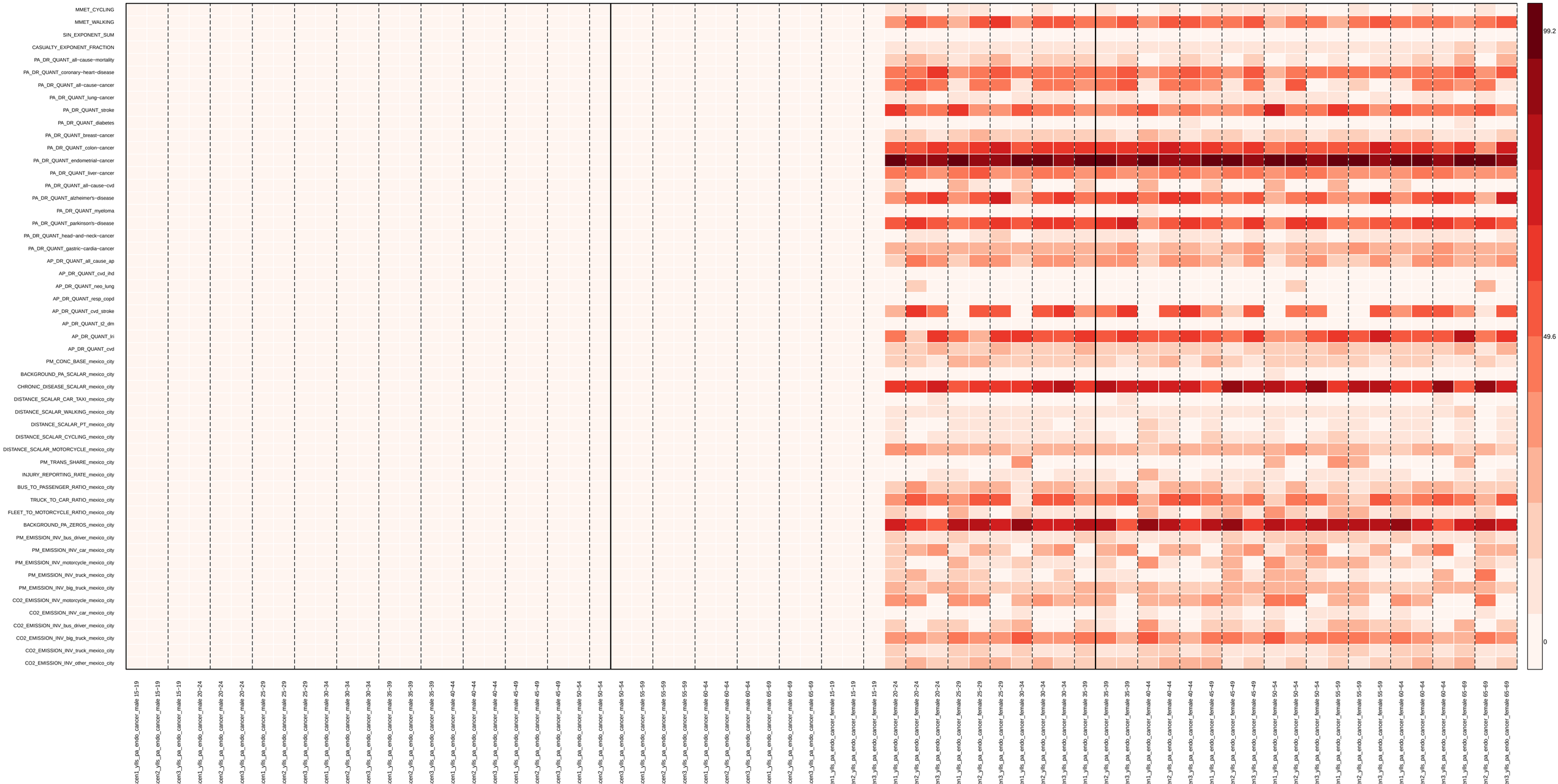
mexico_city − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



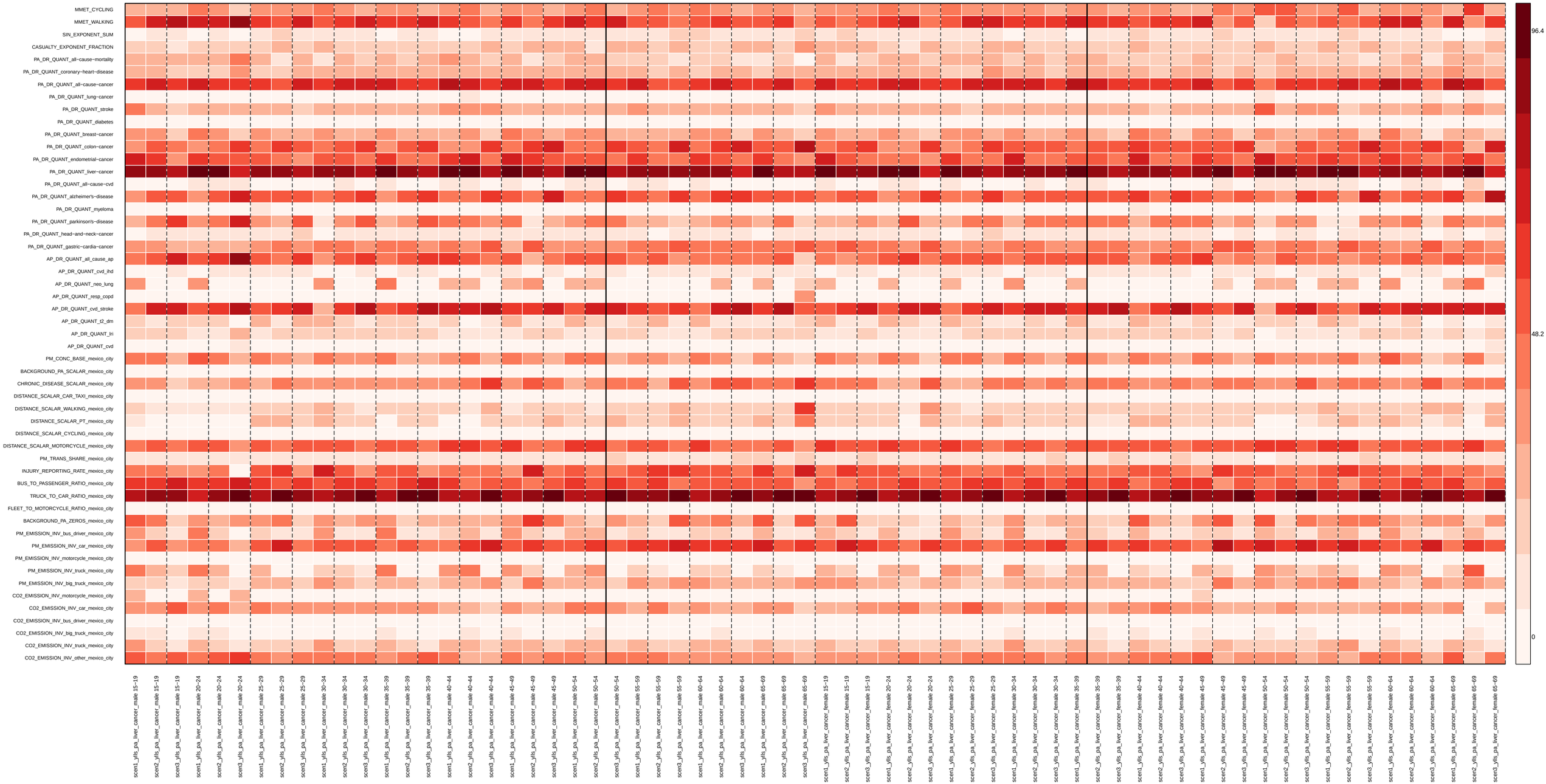
mexico_city − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



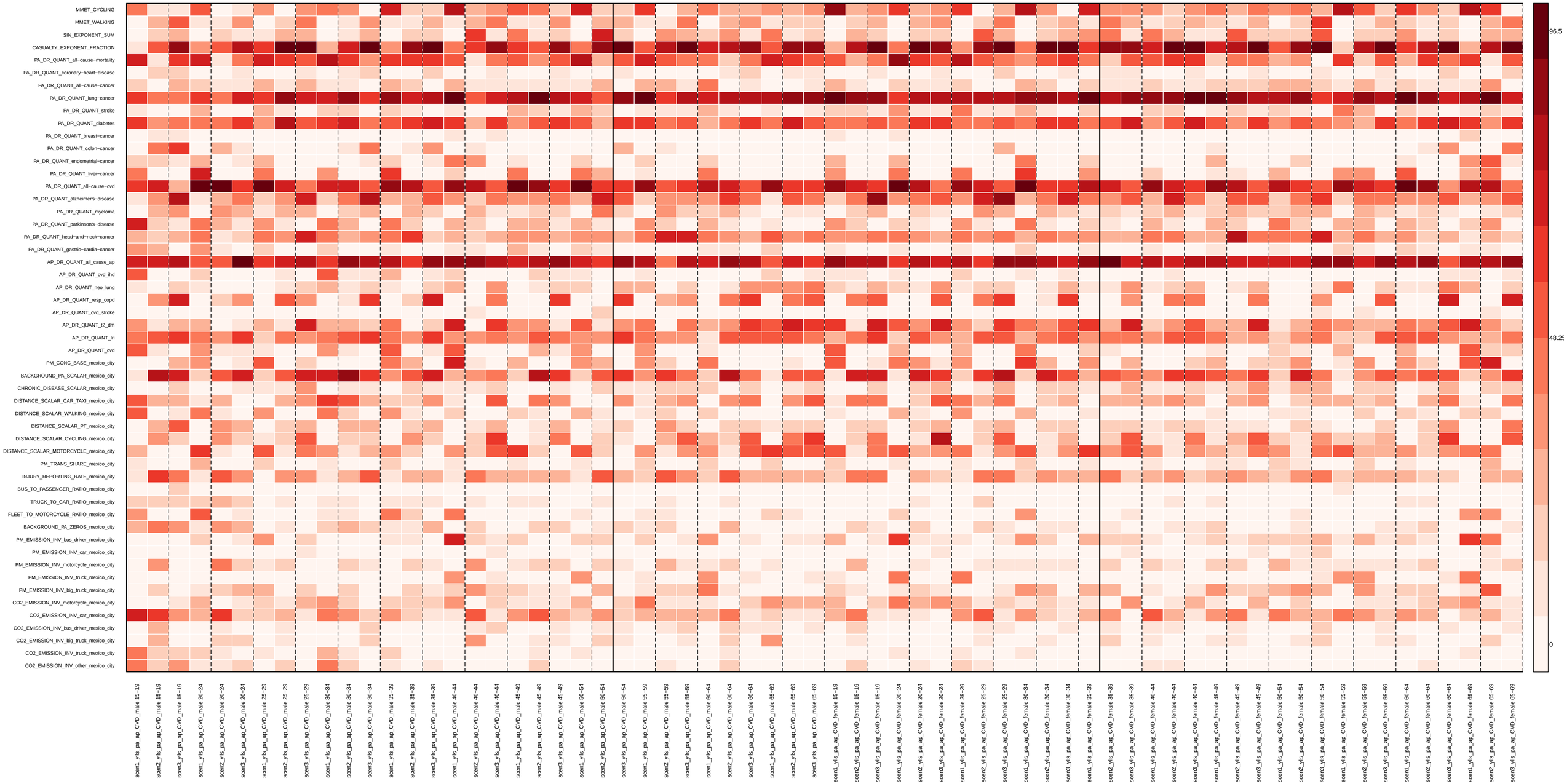
mexico_city − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



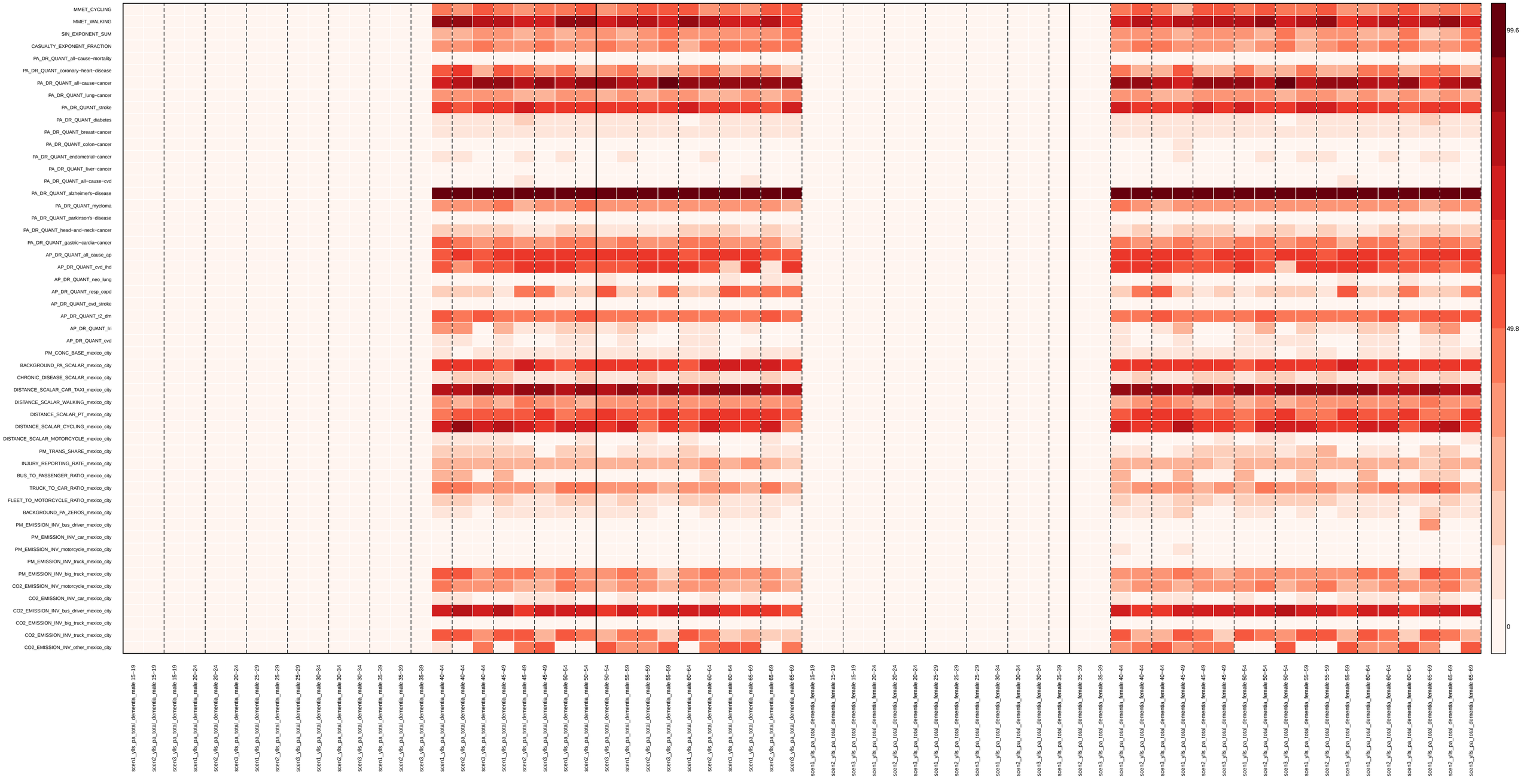
mexico_city − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



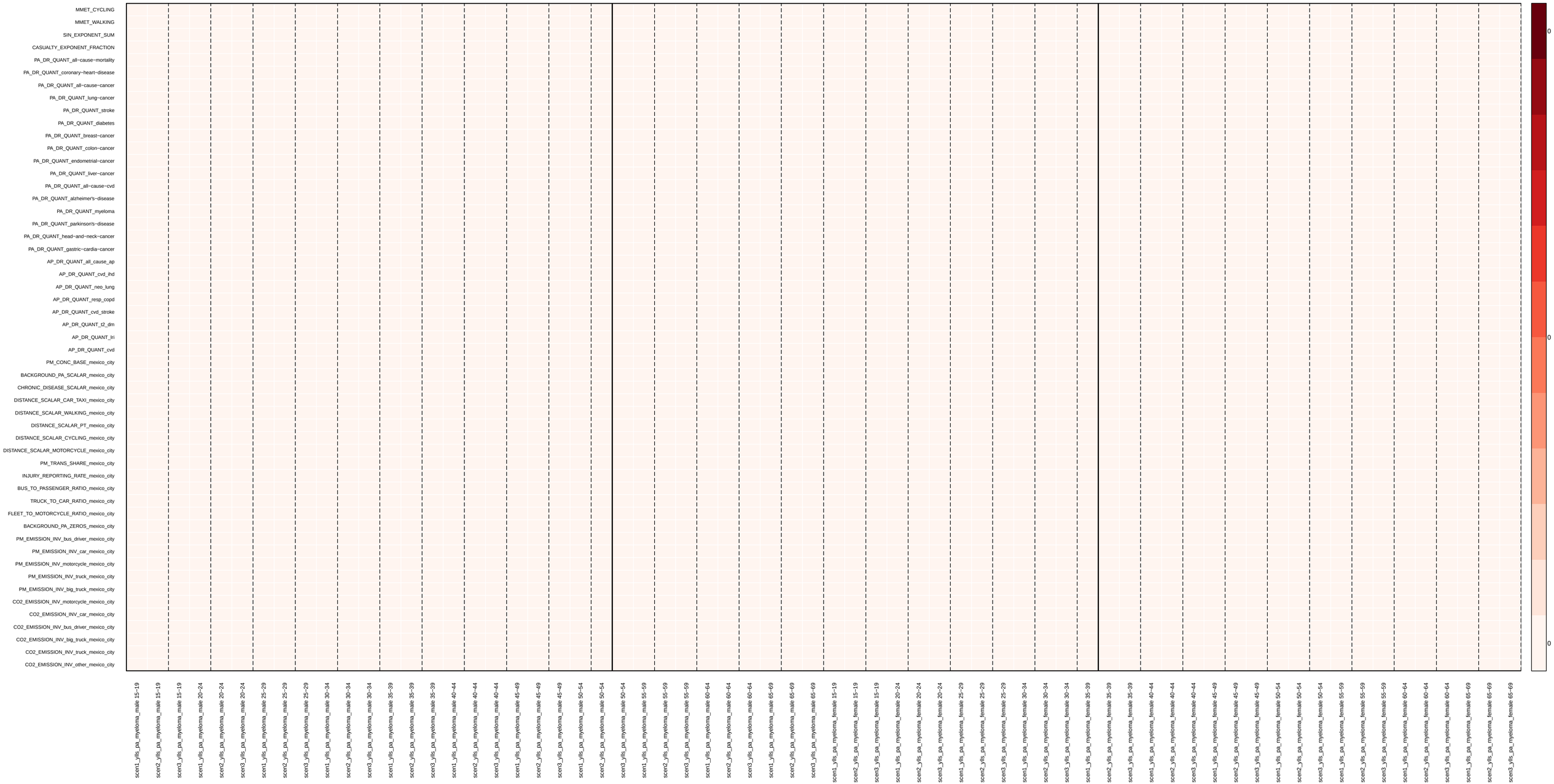
mexico_city − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



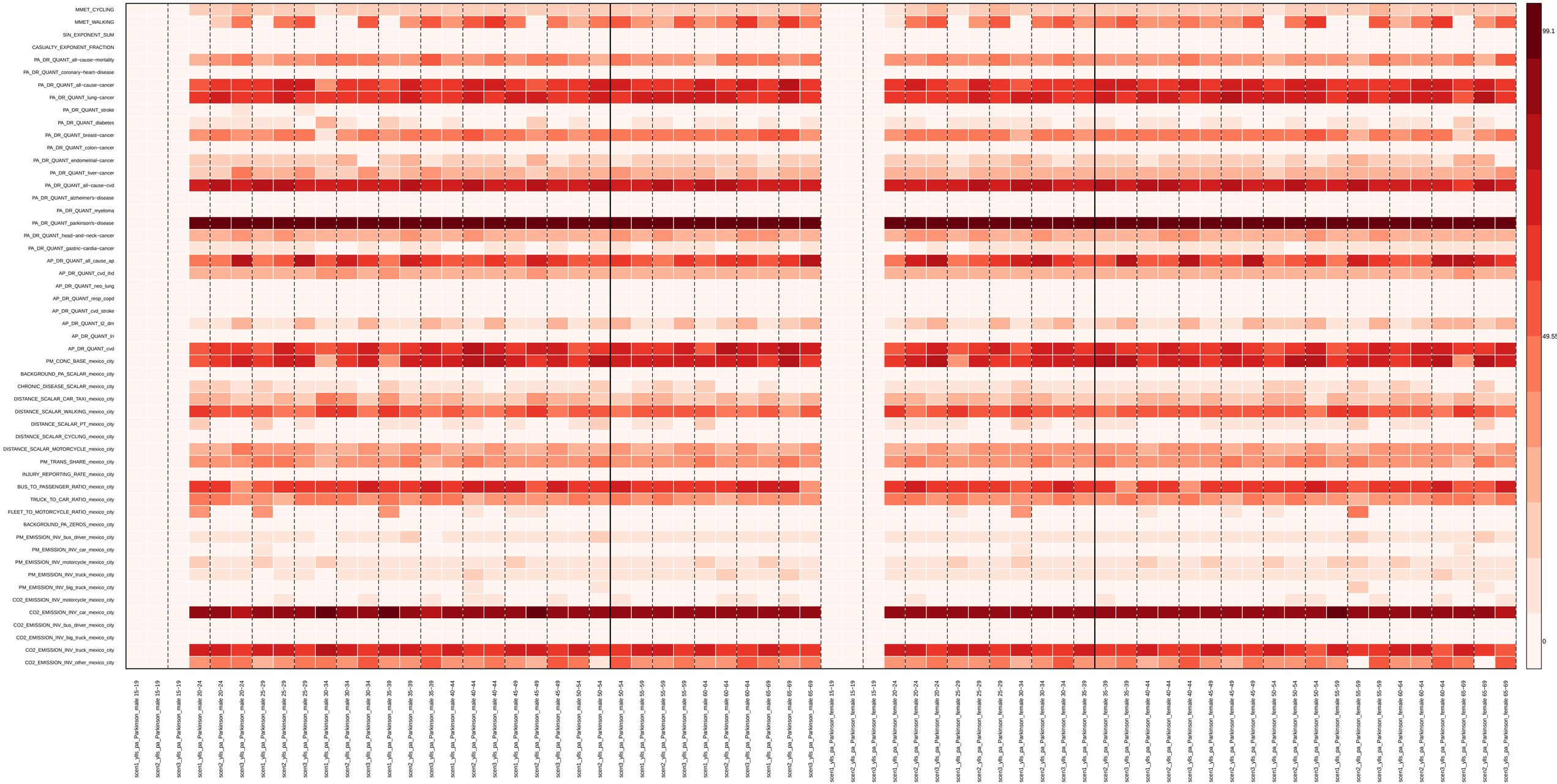
mexico_city − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



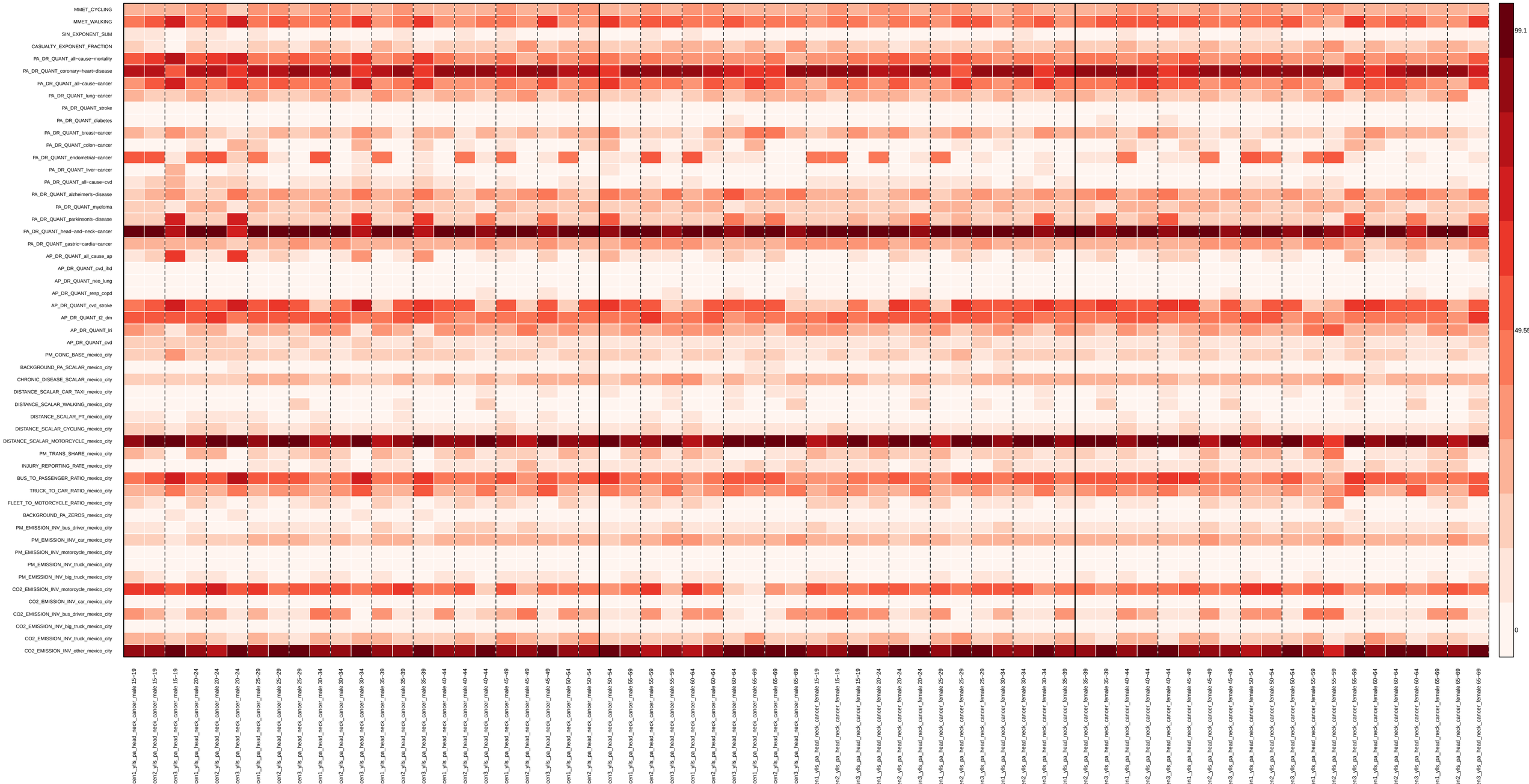
mexico_city − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



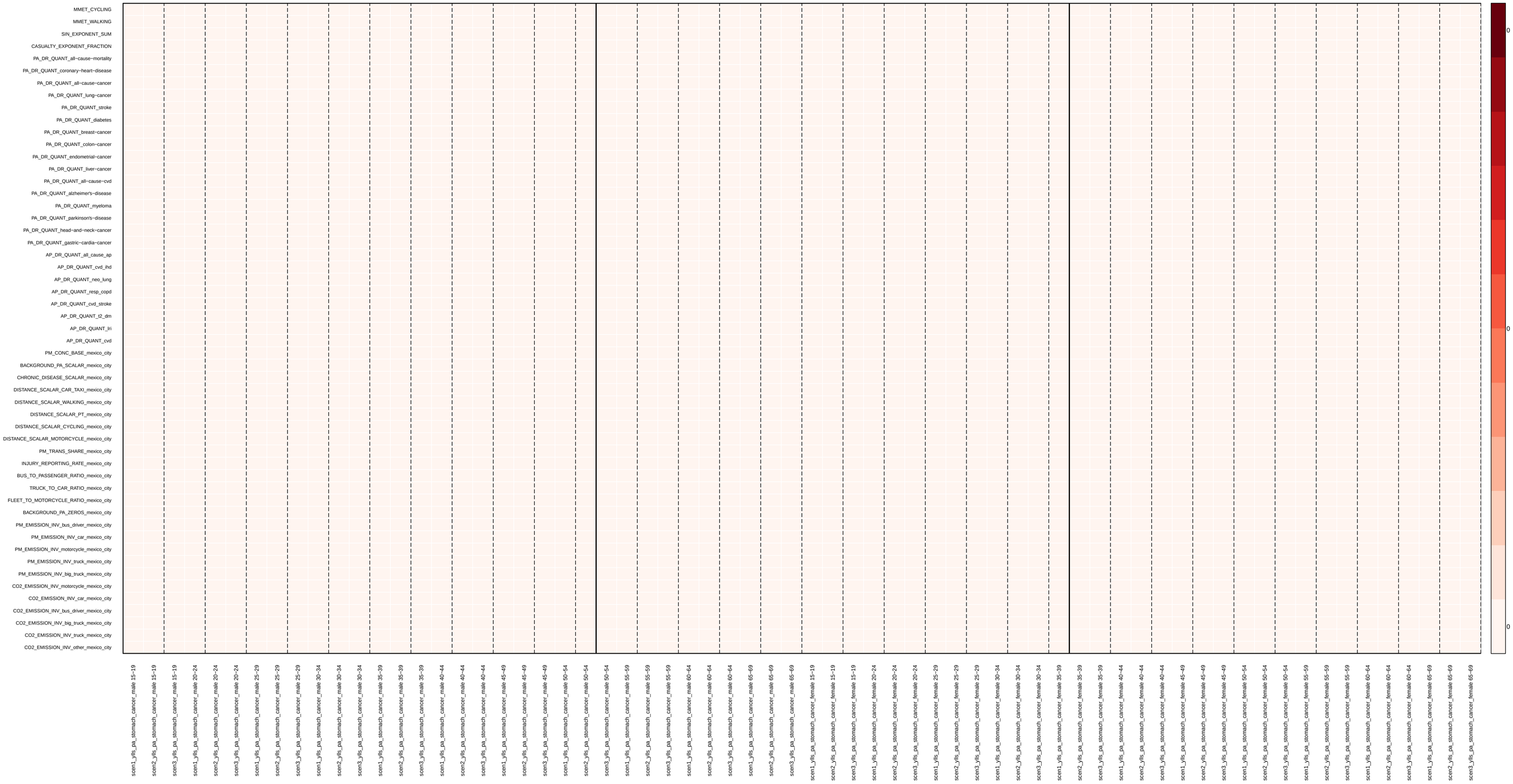
mexico_city − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



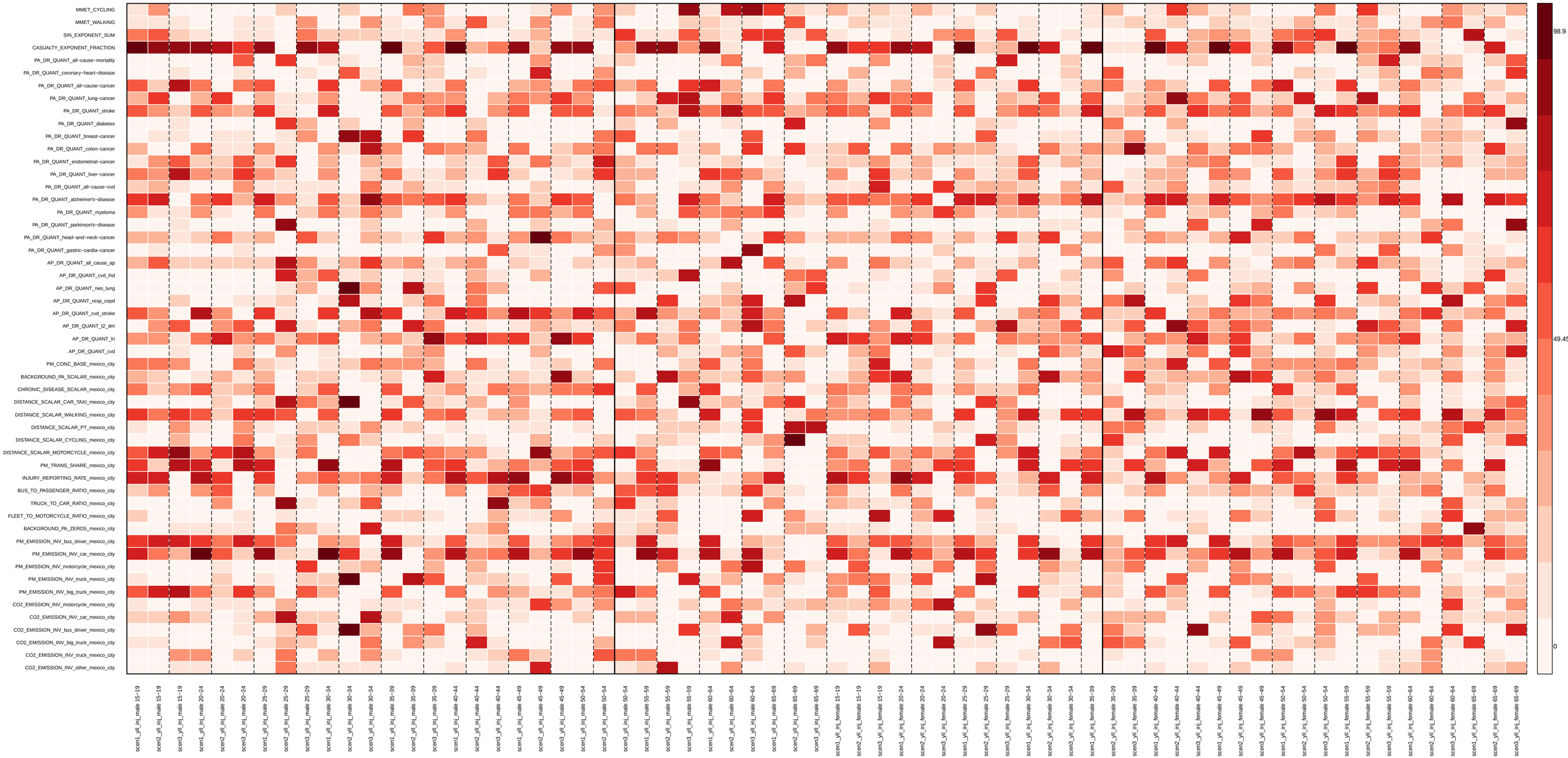
mexico_city − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



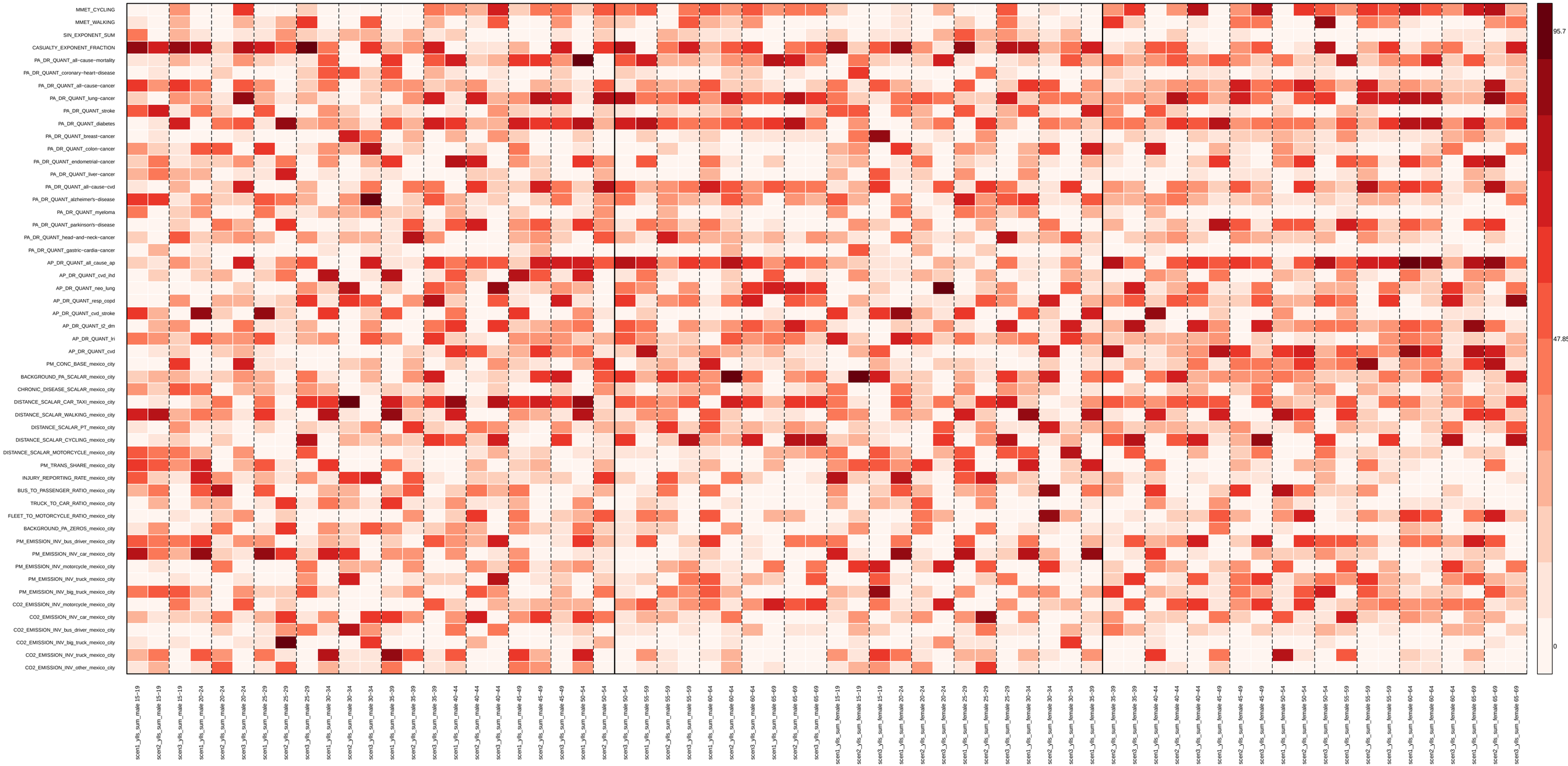
mexico_city − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



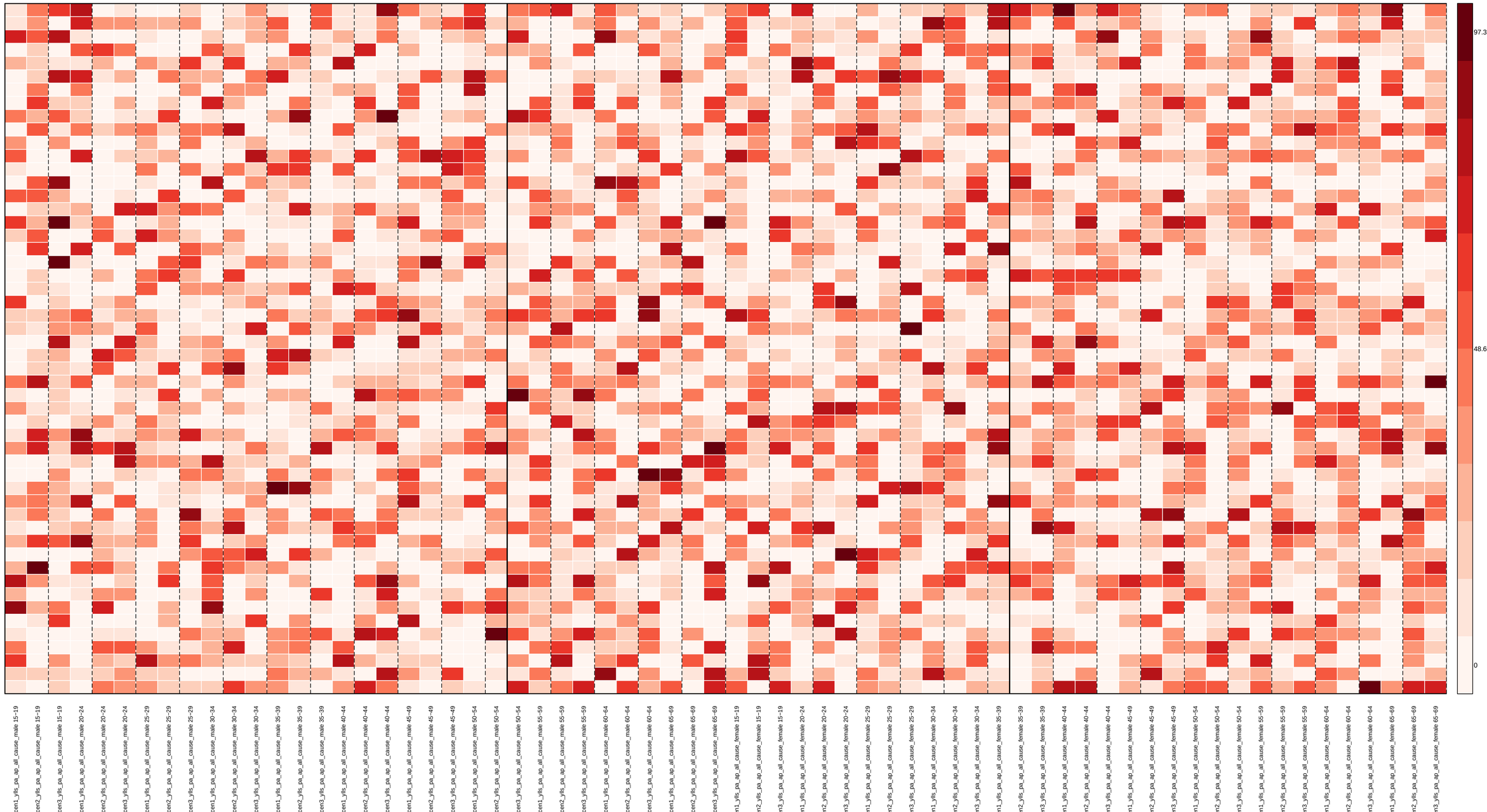
mexico_city − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



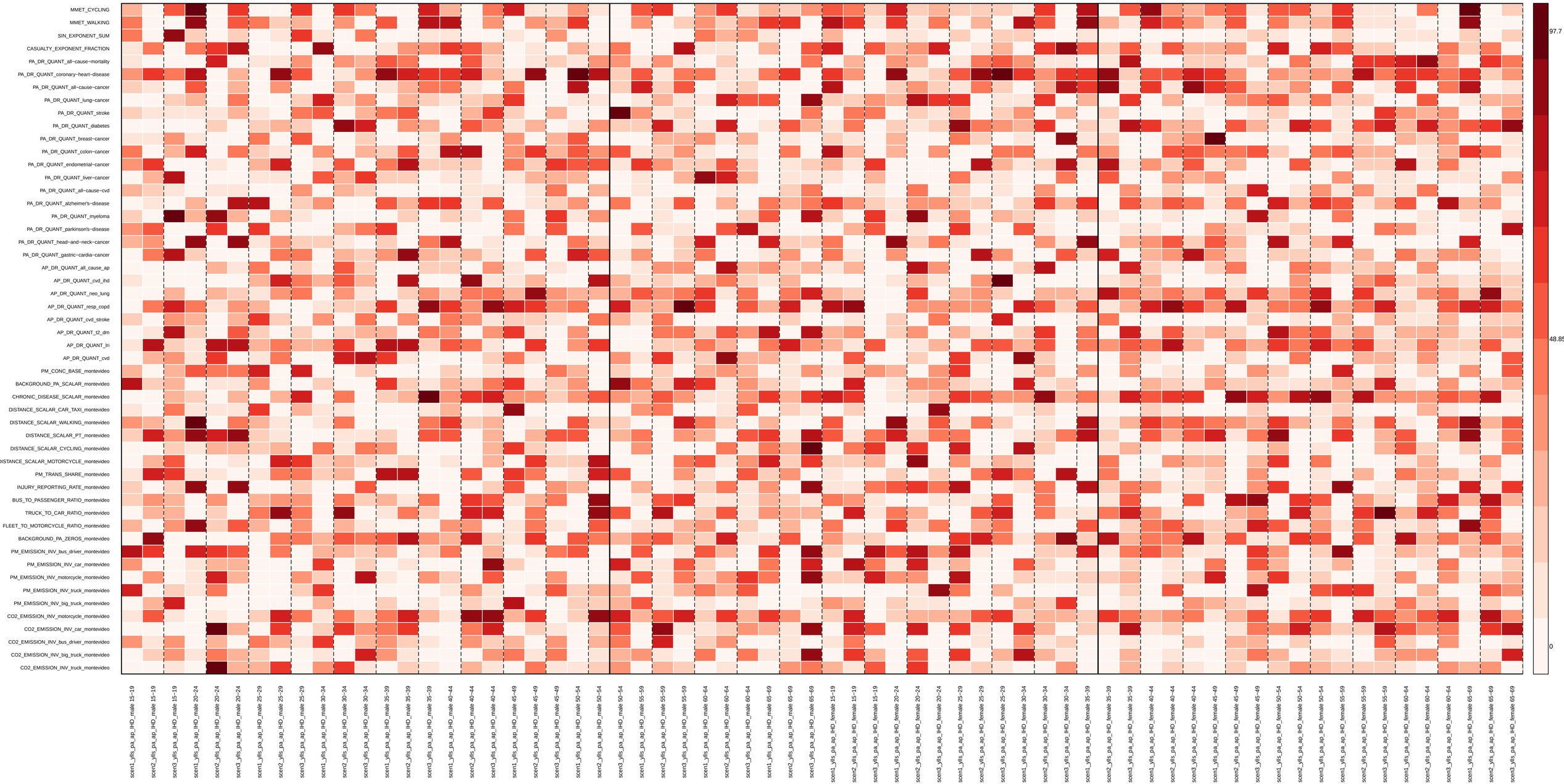
mexico_city − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



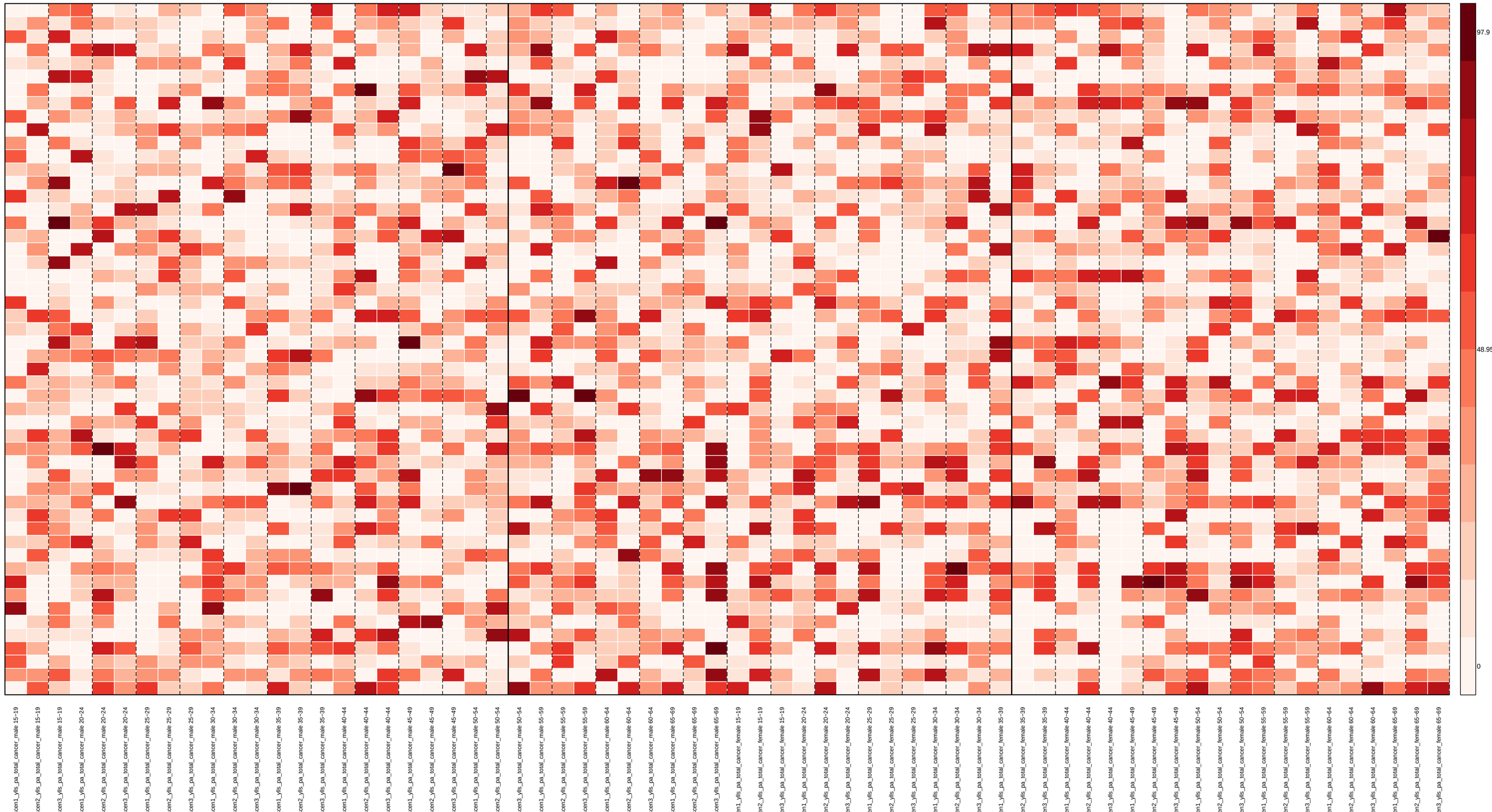
montevideo − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



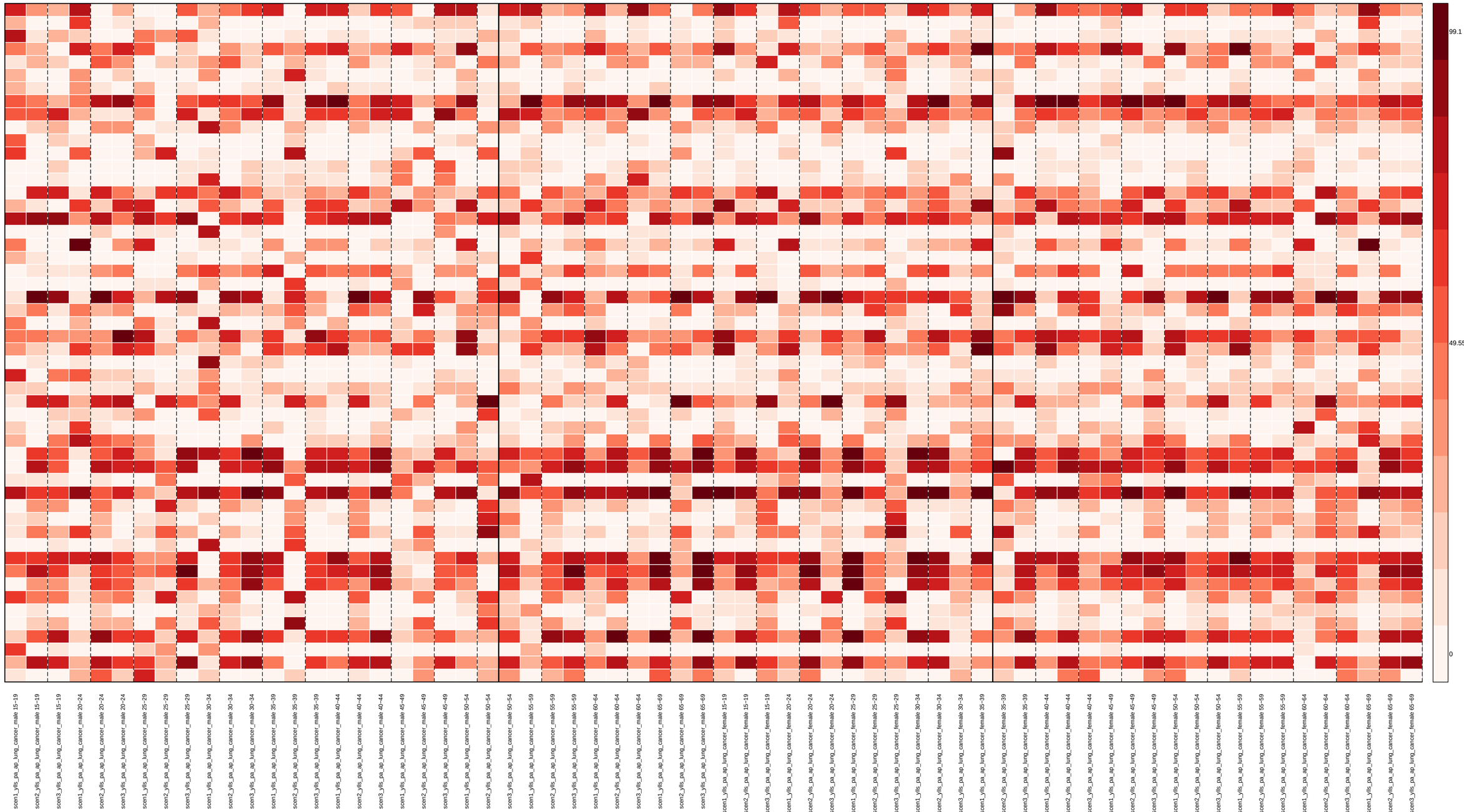
montevideo − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



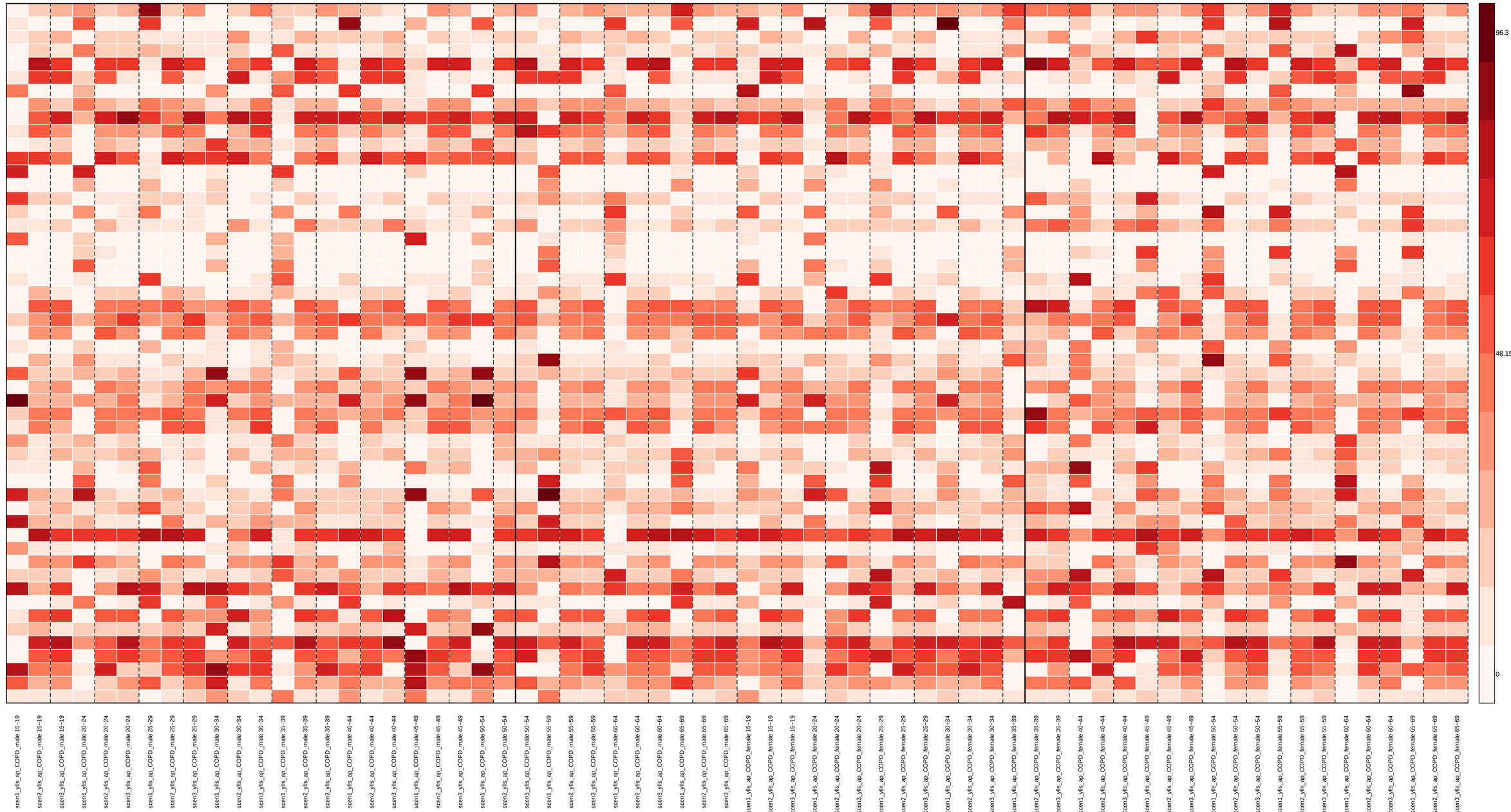
montevideo − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



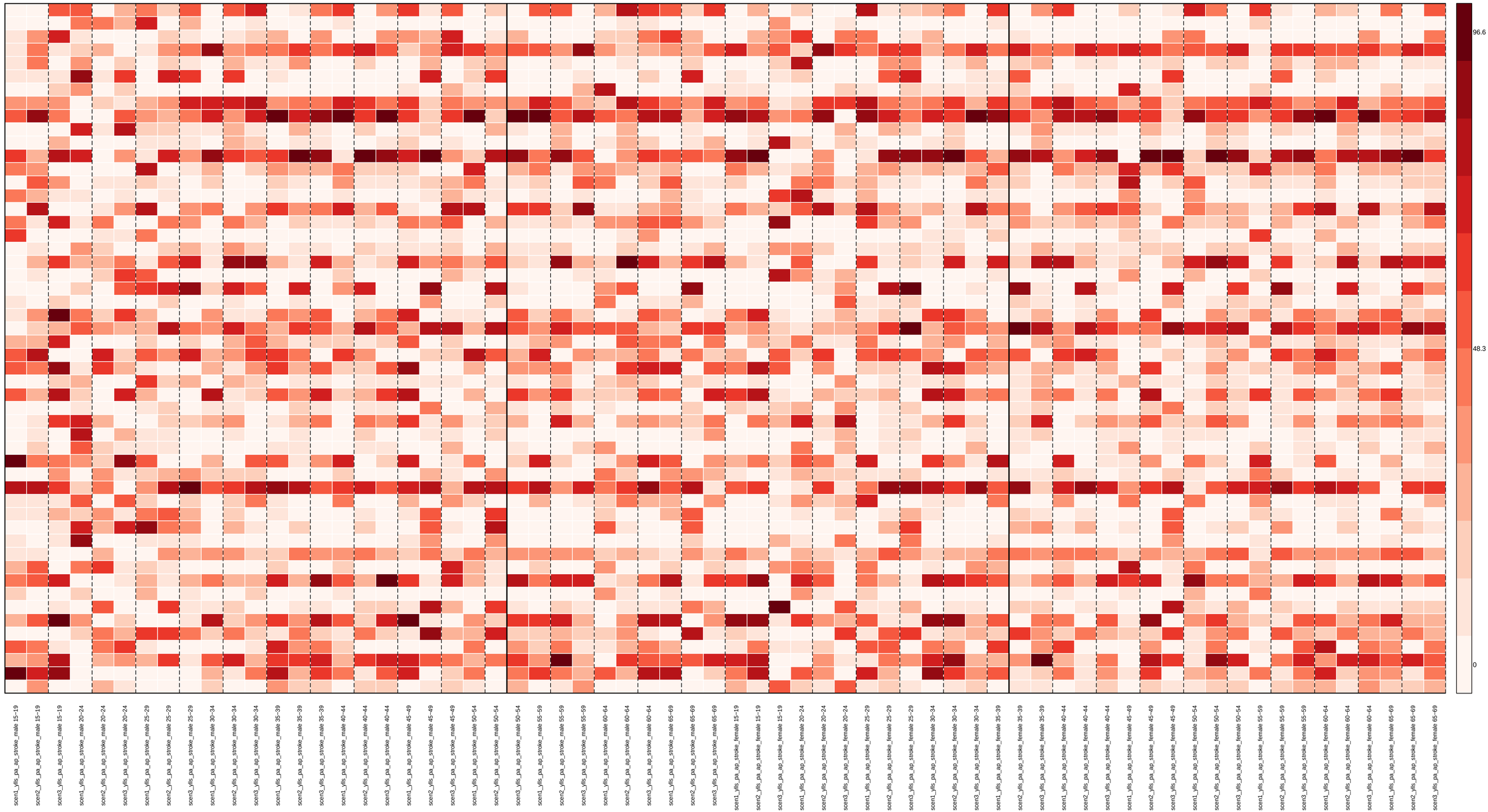
montevideo − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



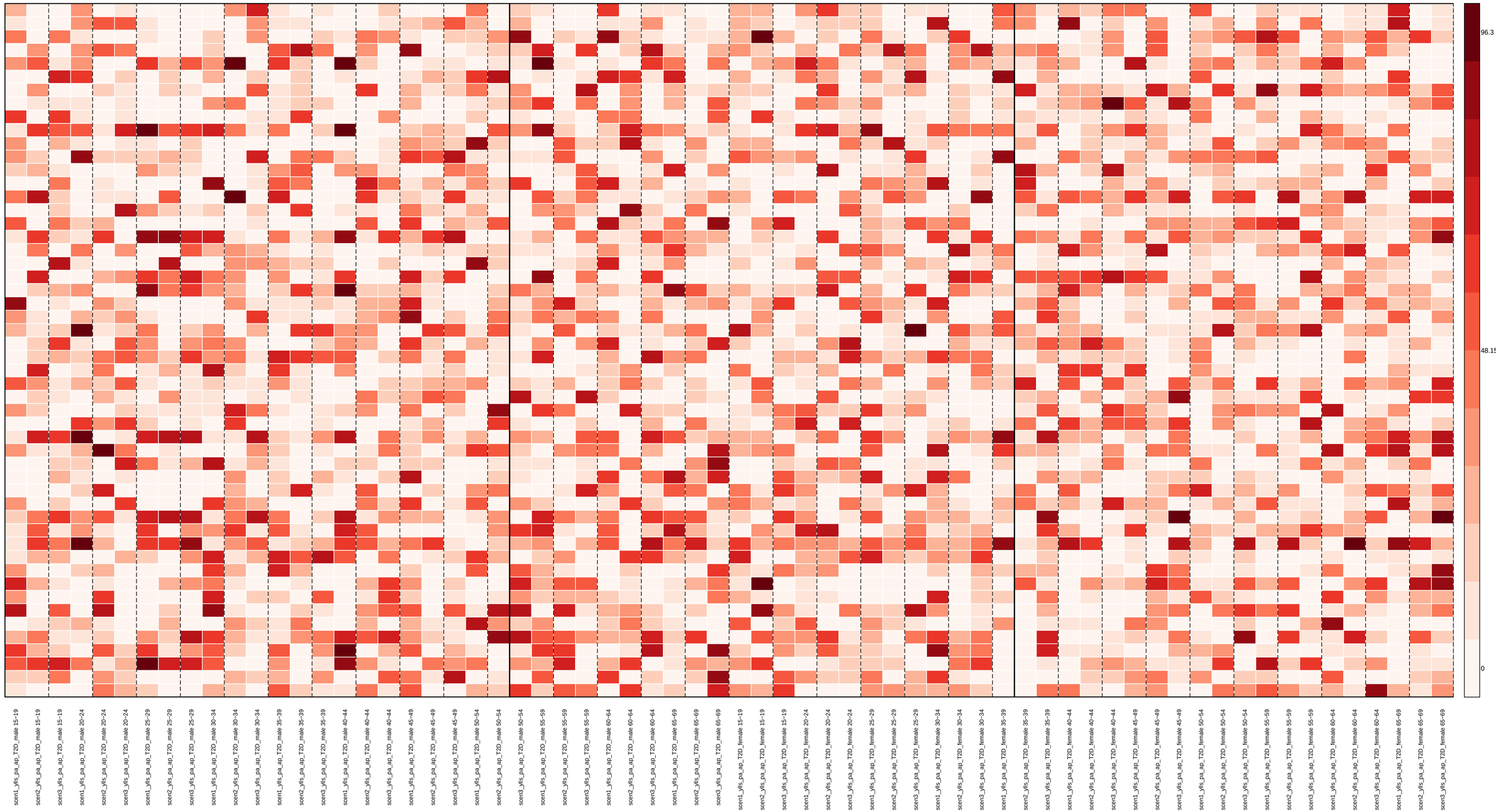
montevideo − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



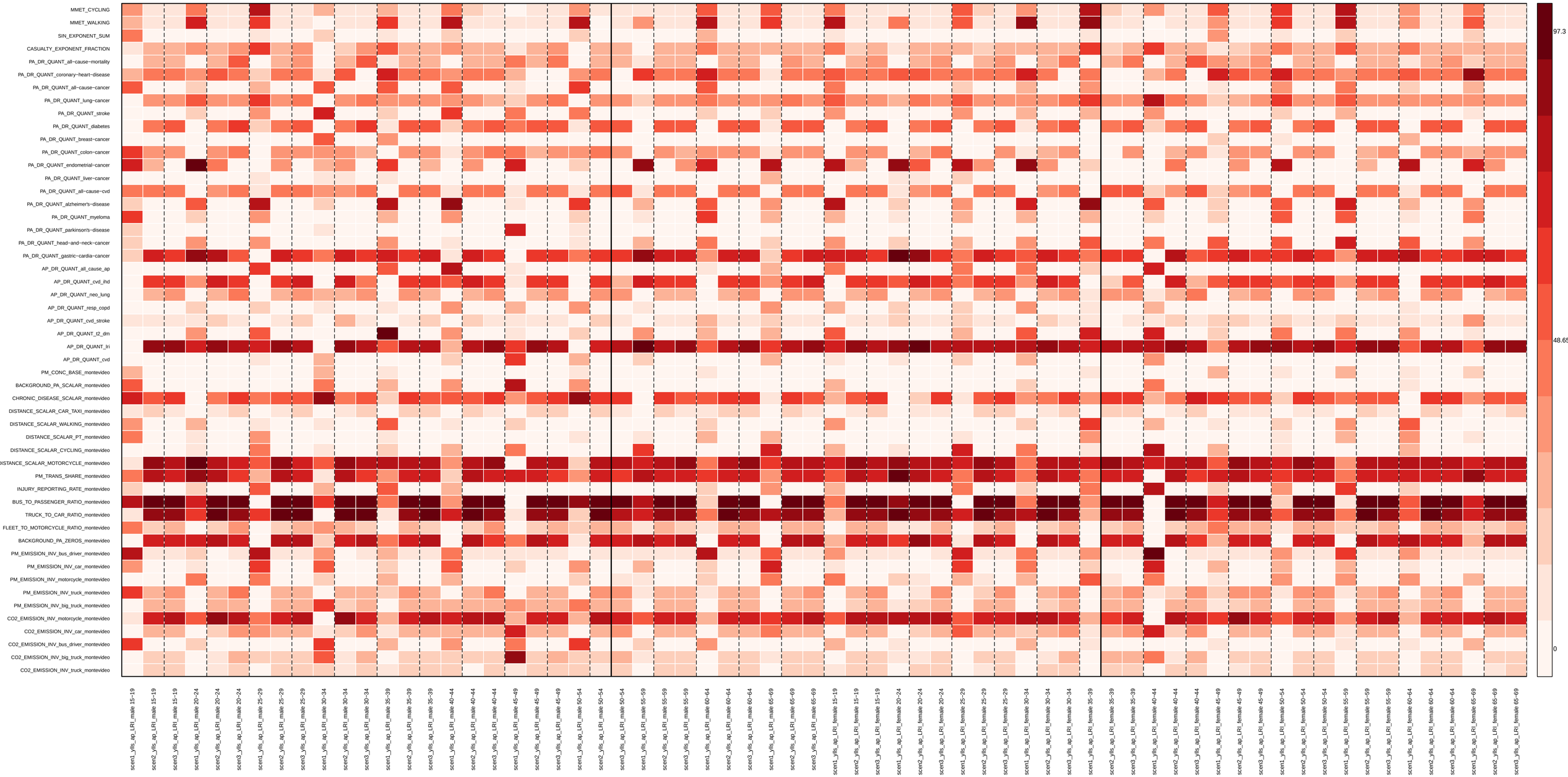
montevideo − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



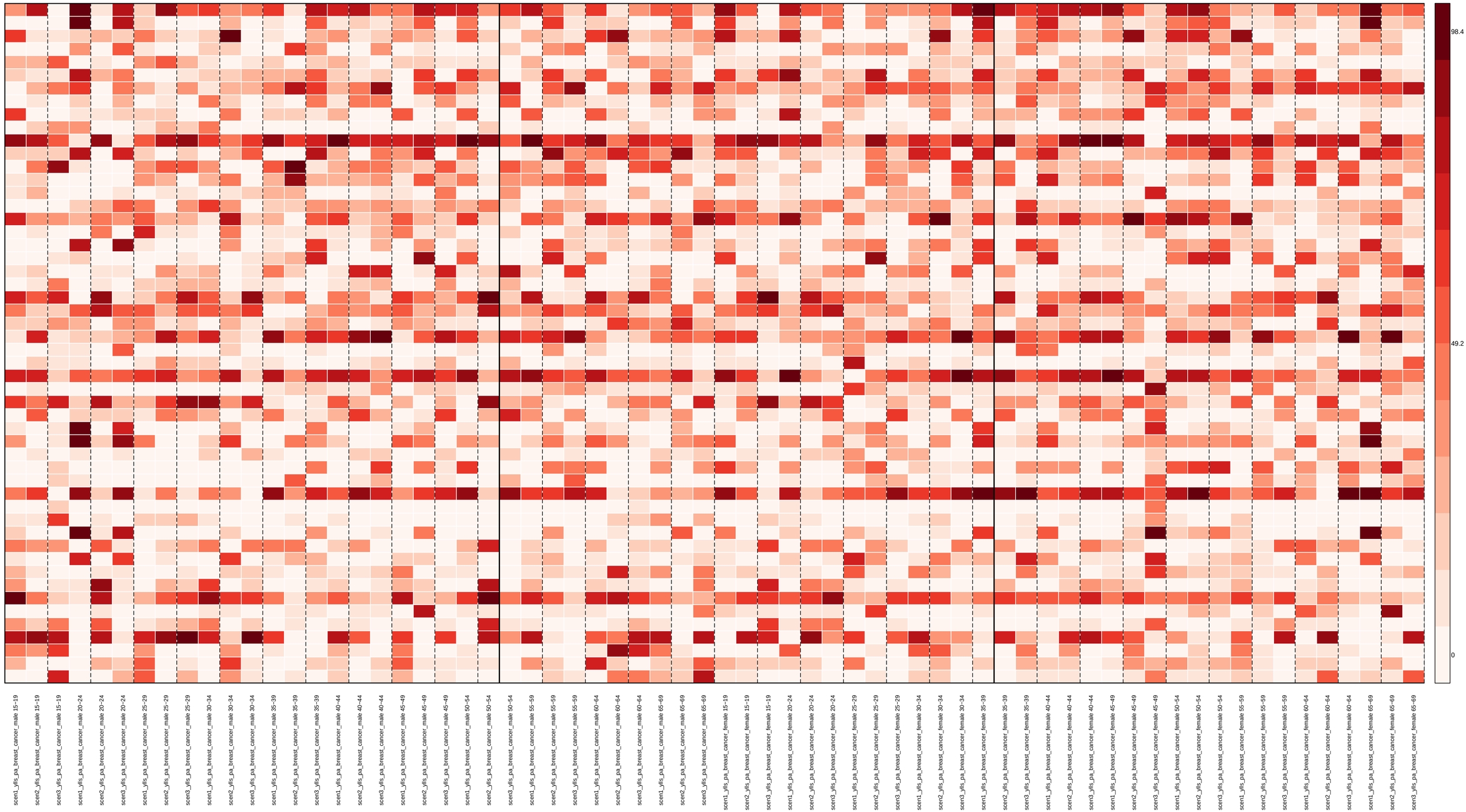
montevideo − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



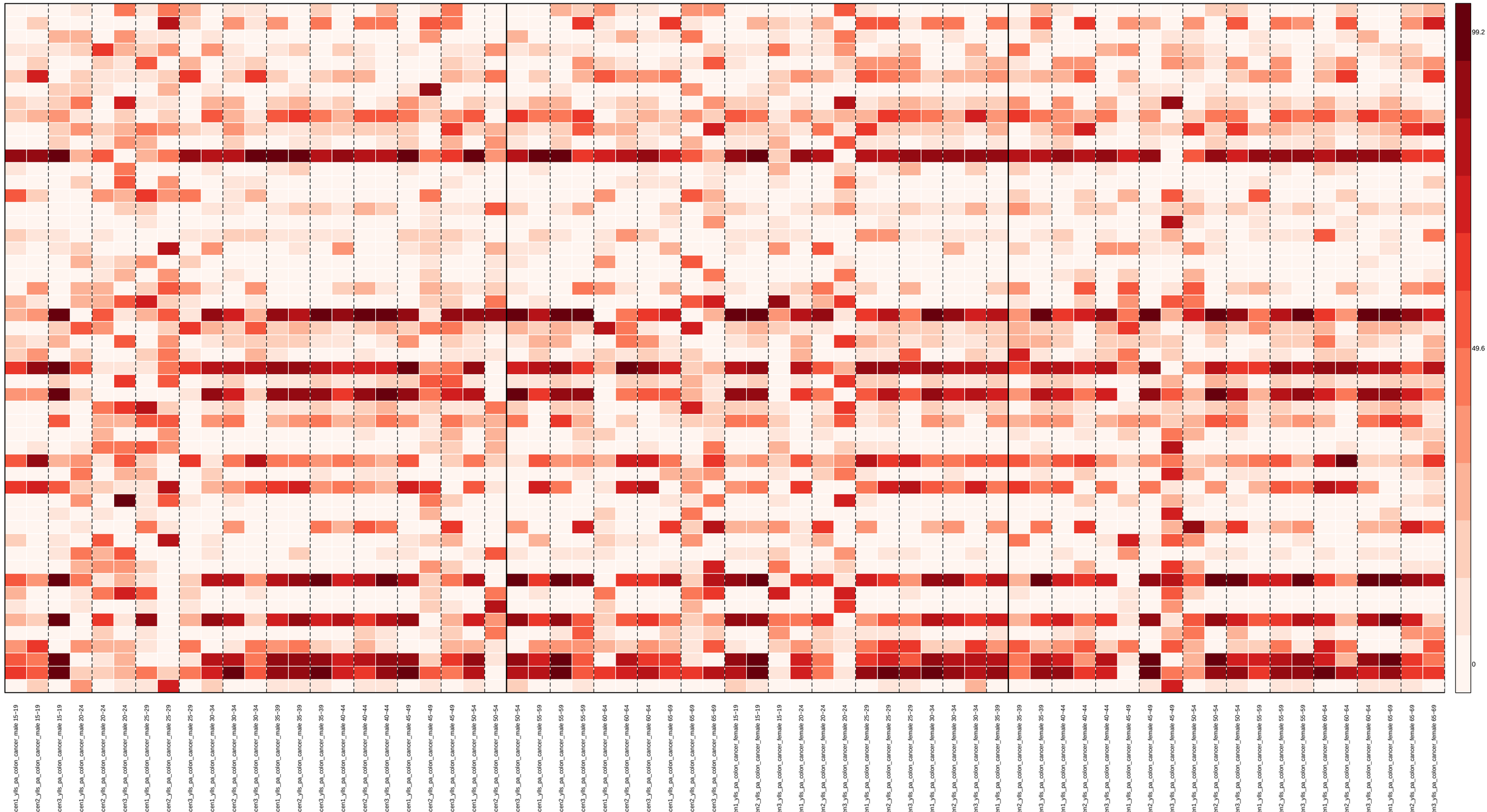
montevideo − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



montevideo − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

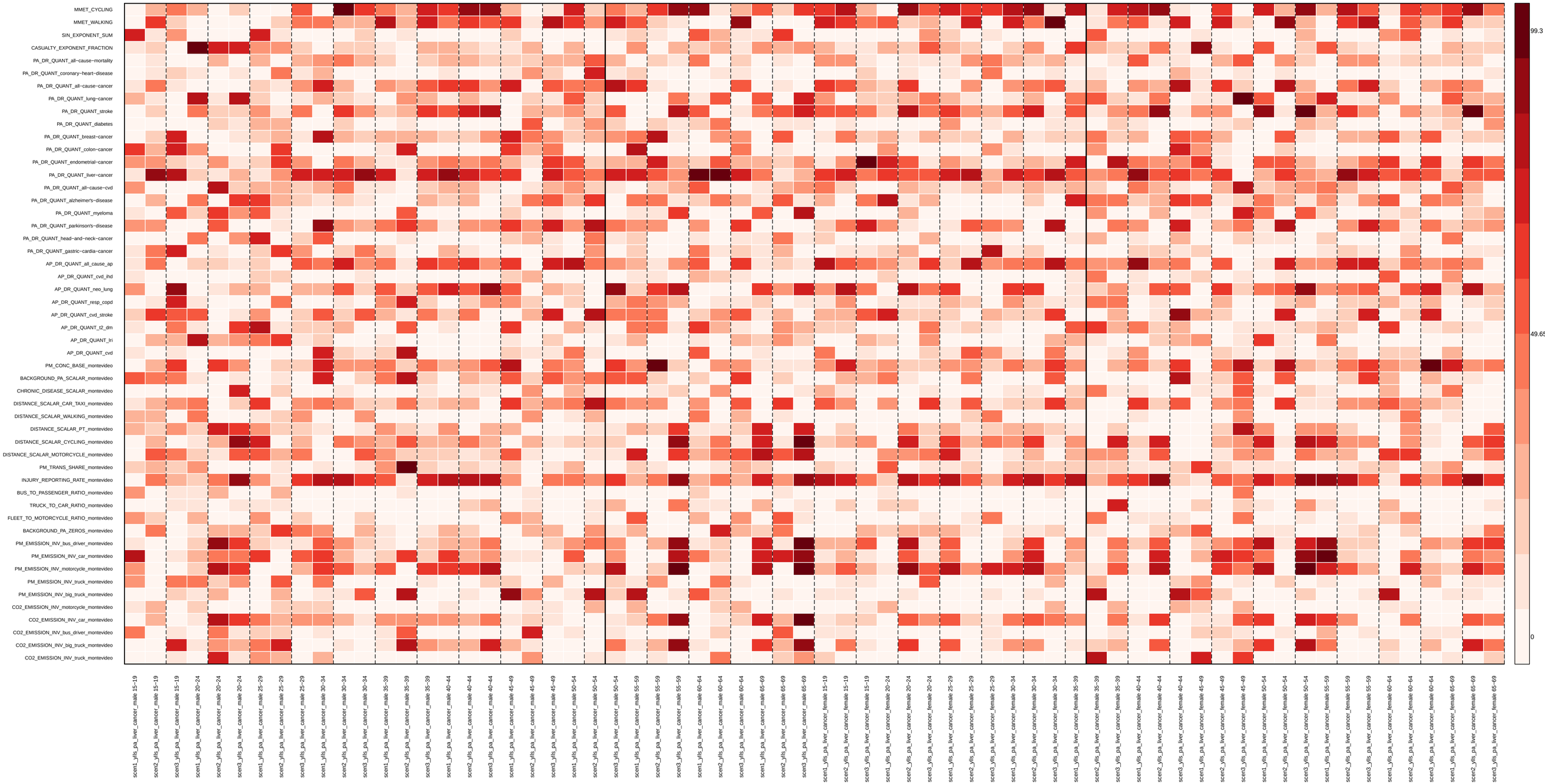


montevideo − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

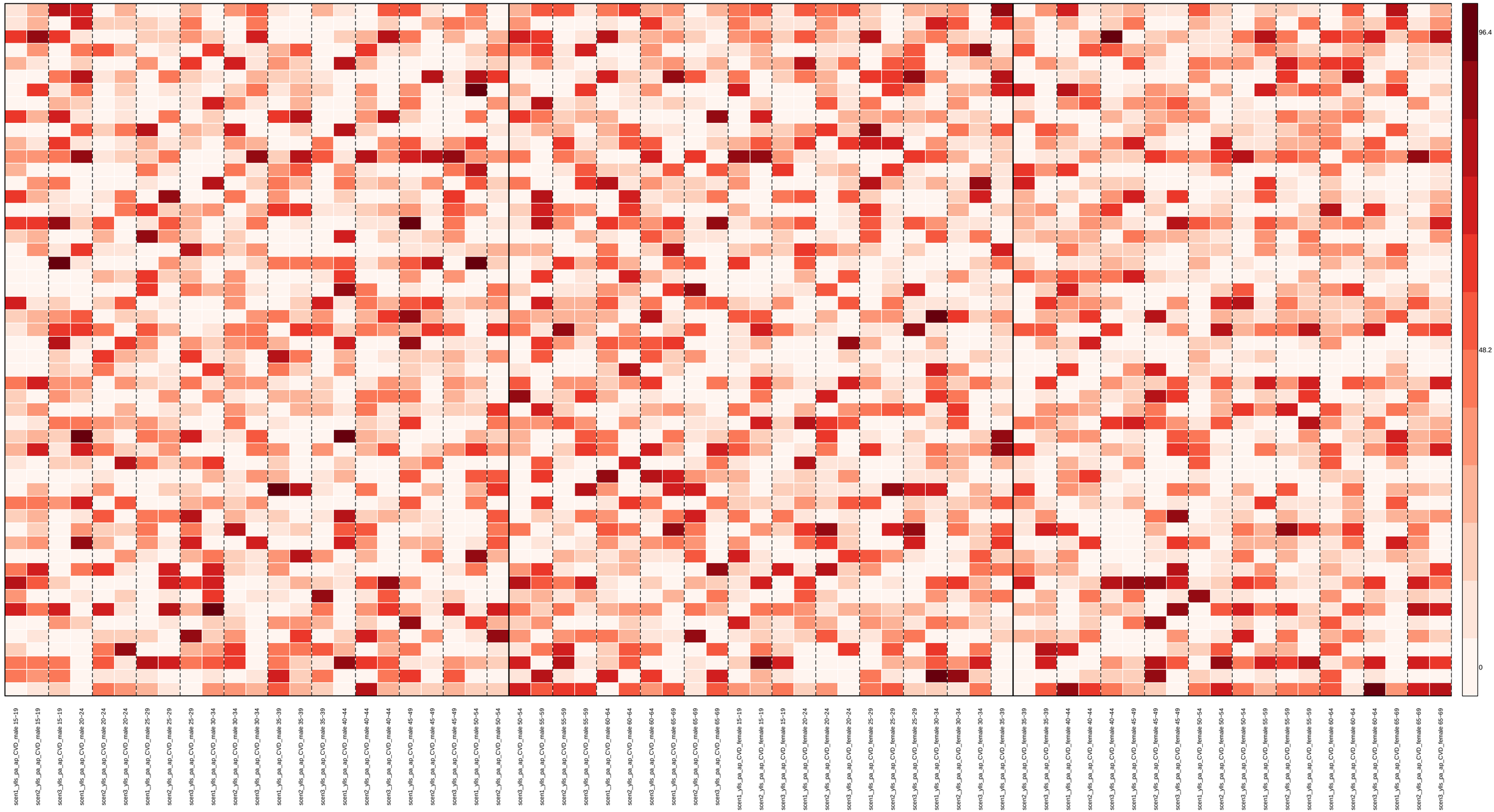


montevideo − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

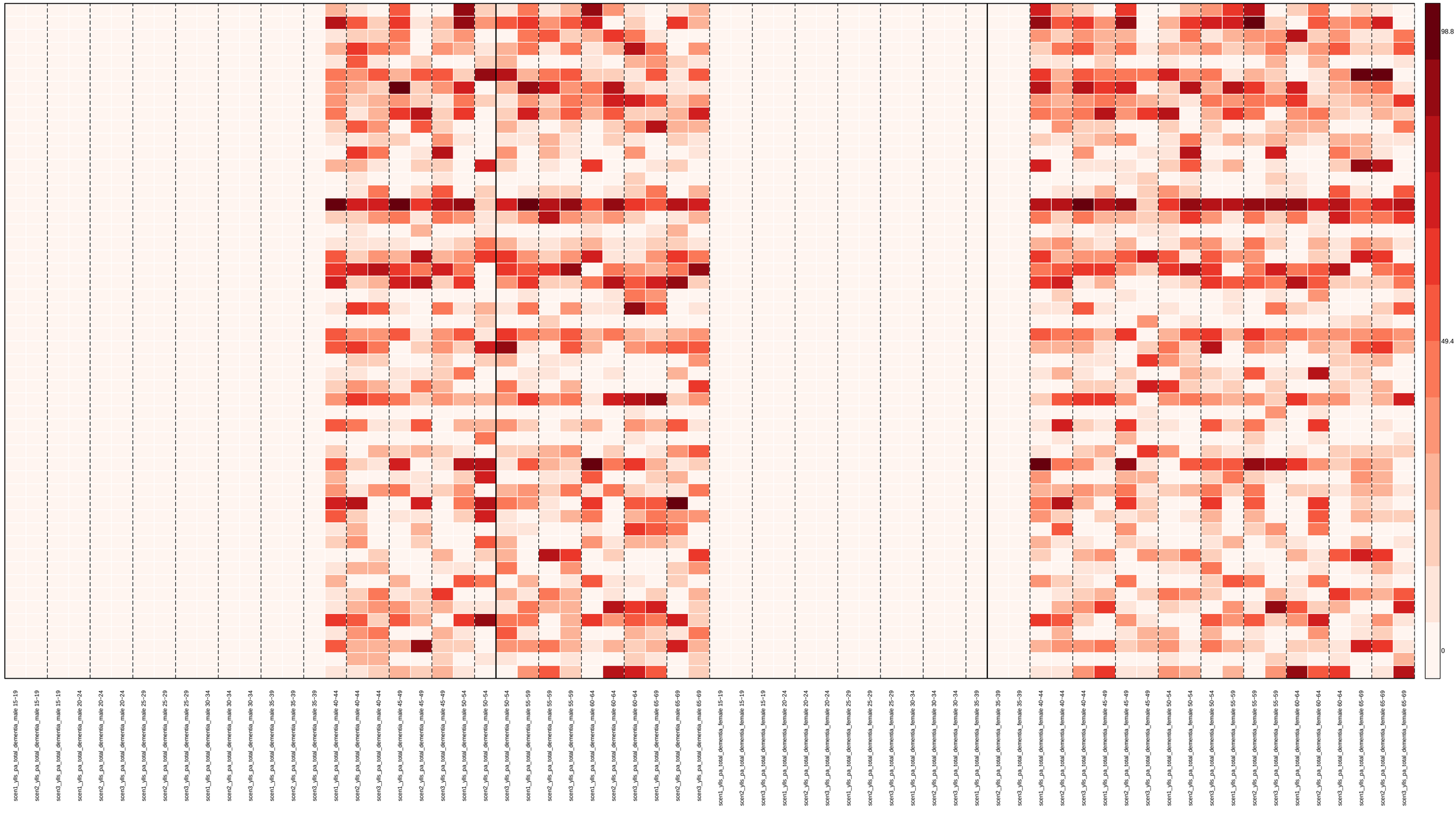
montevideo − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



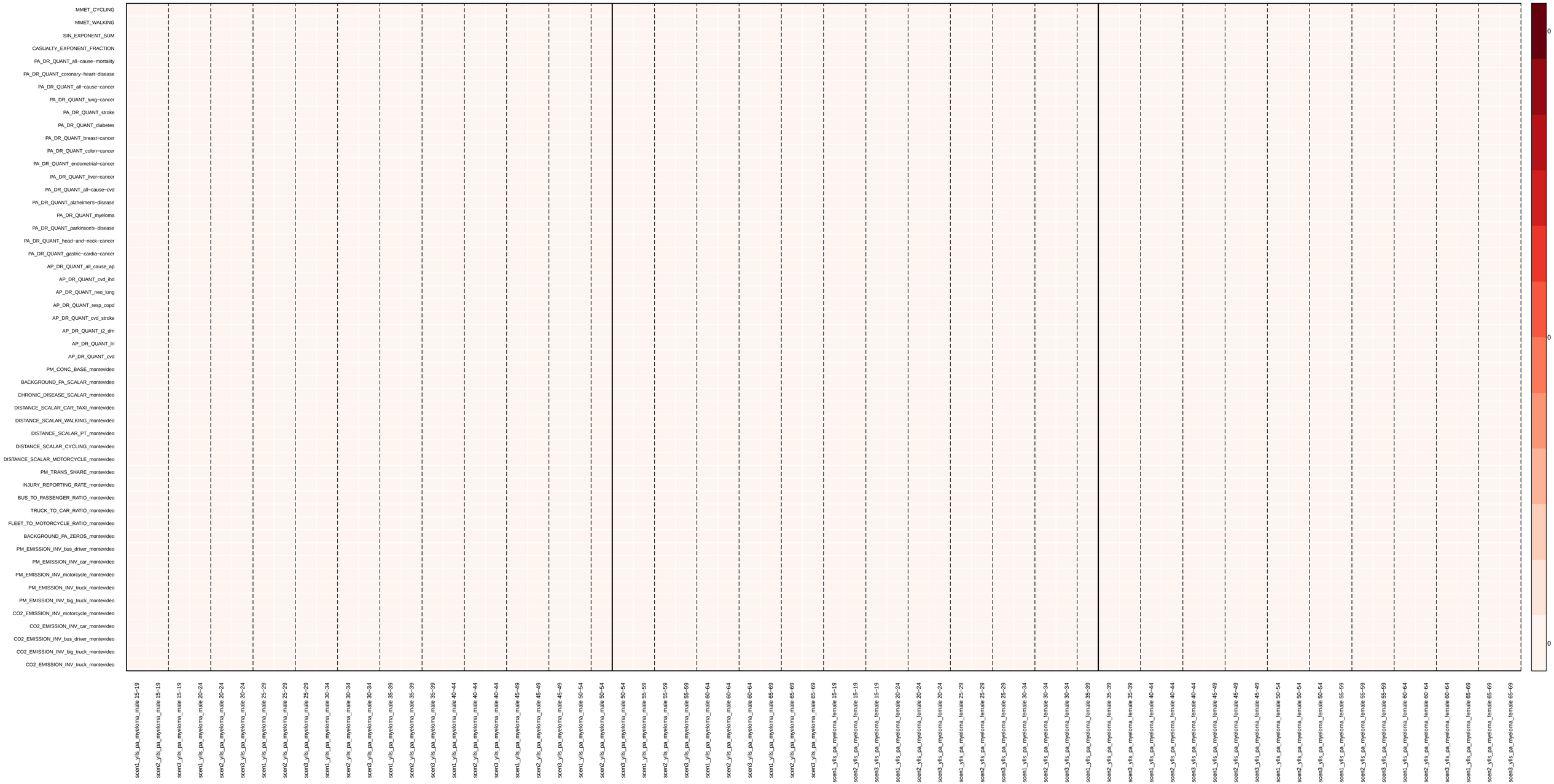
montevideo − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



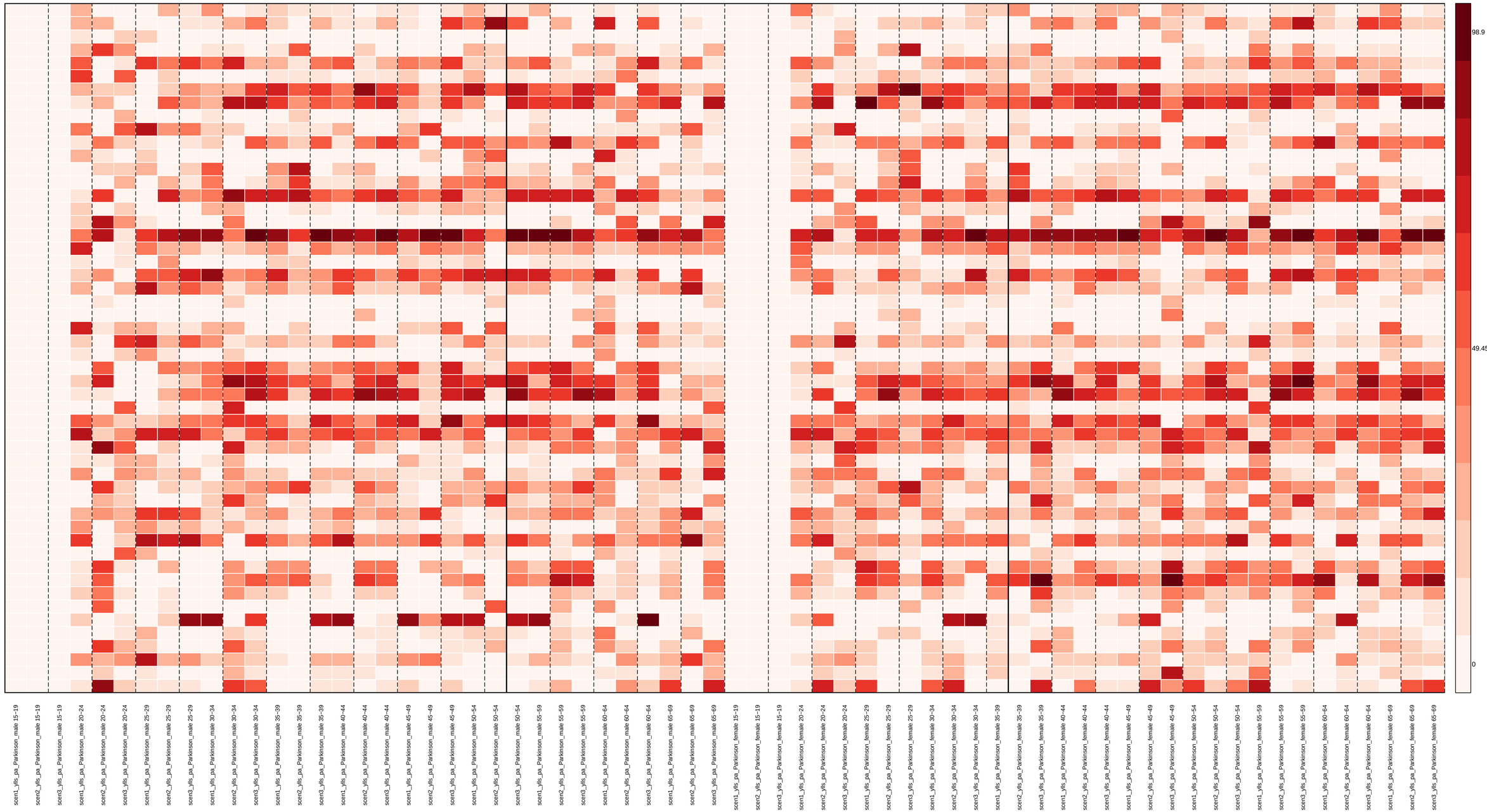
montevideo − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



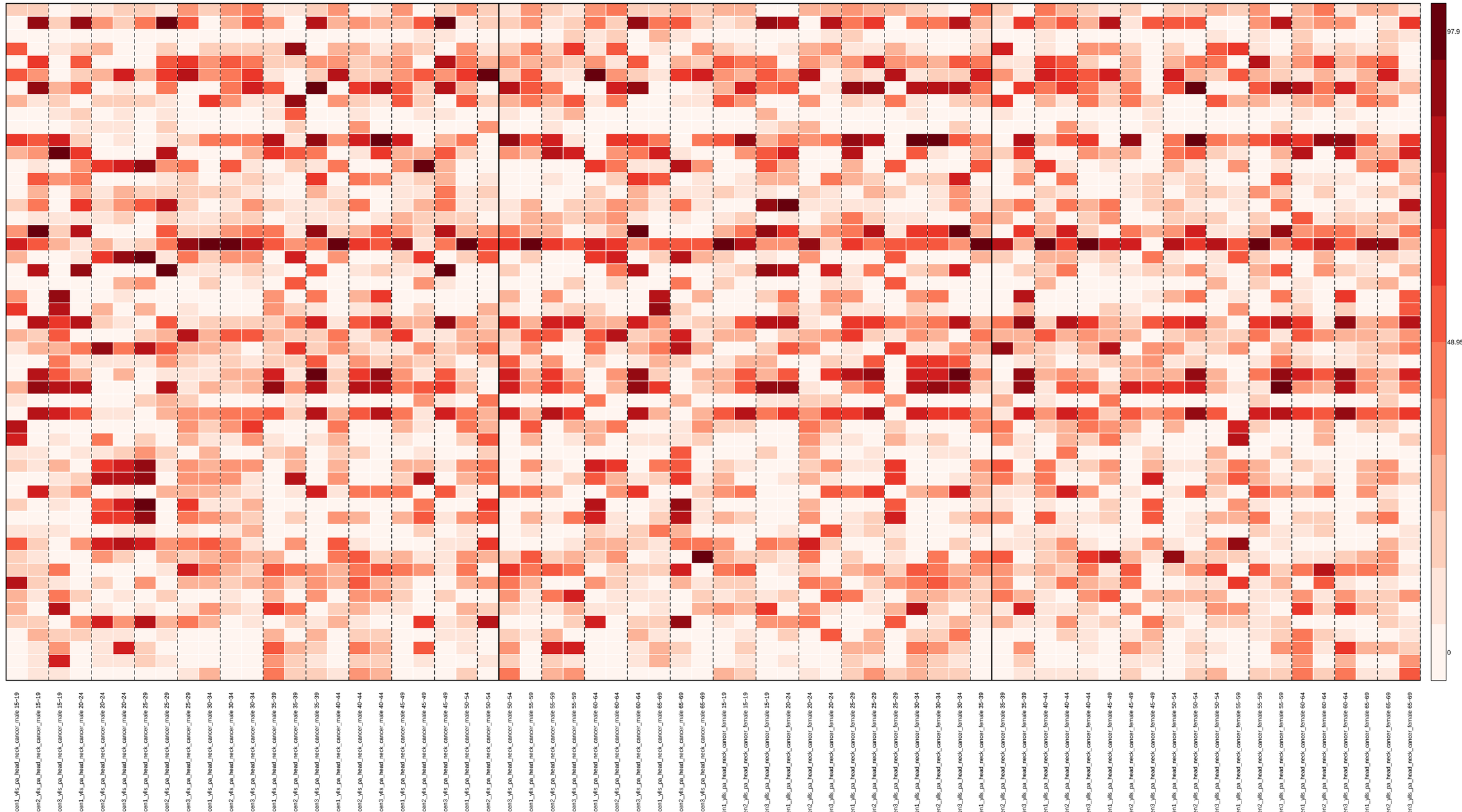
montevideo − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



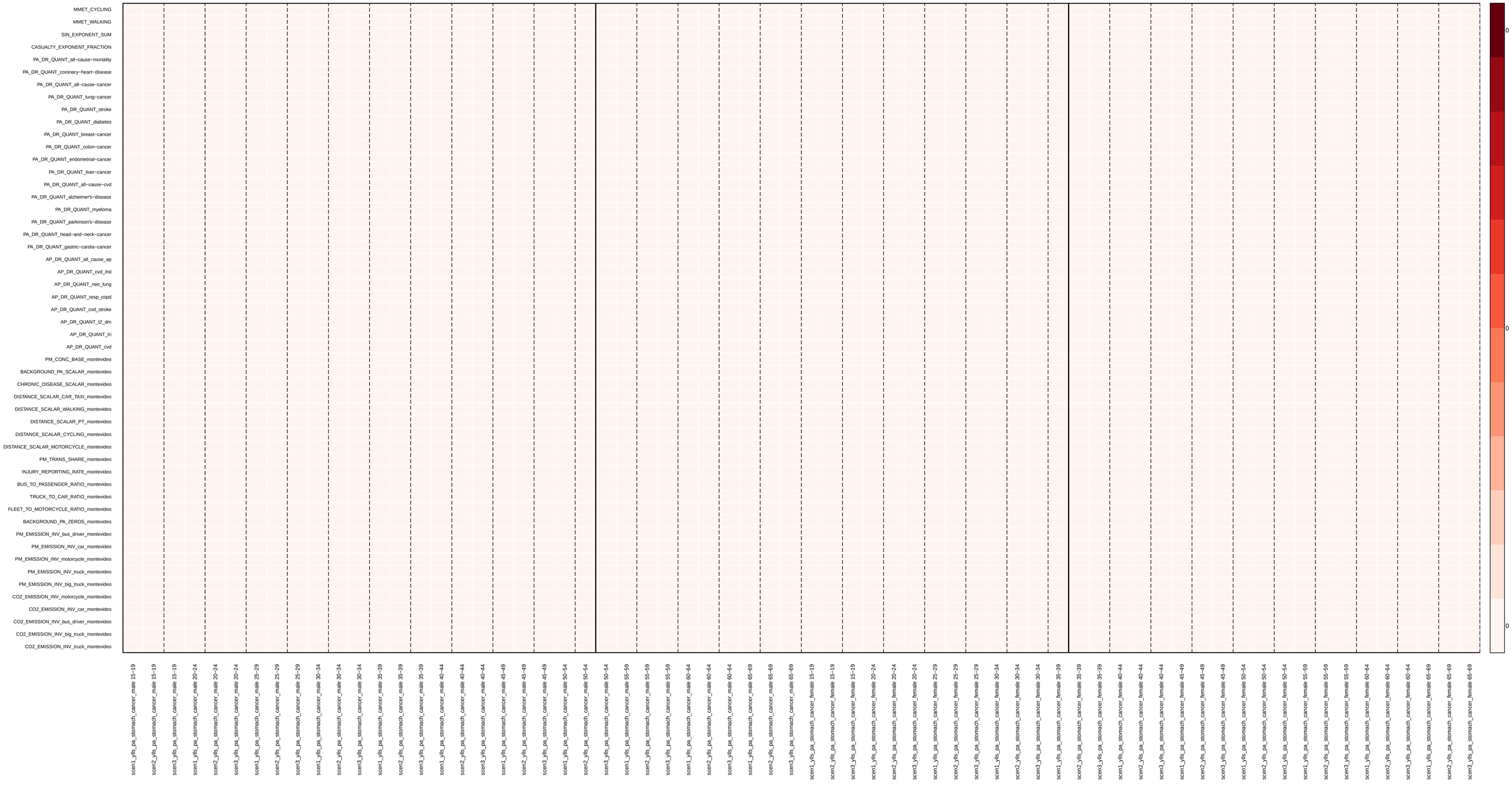
montevideo − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



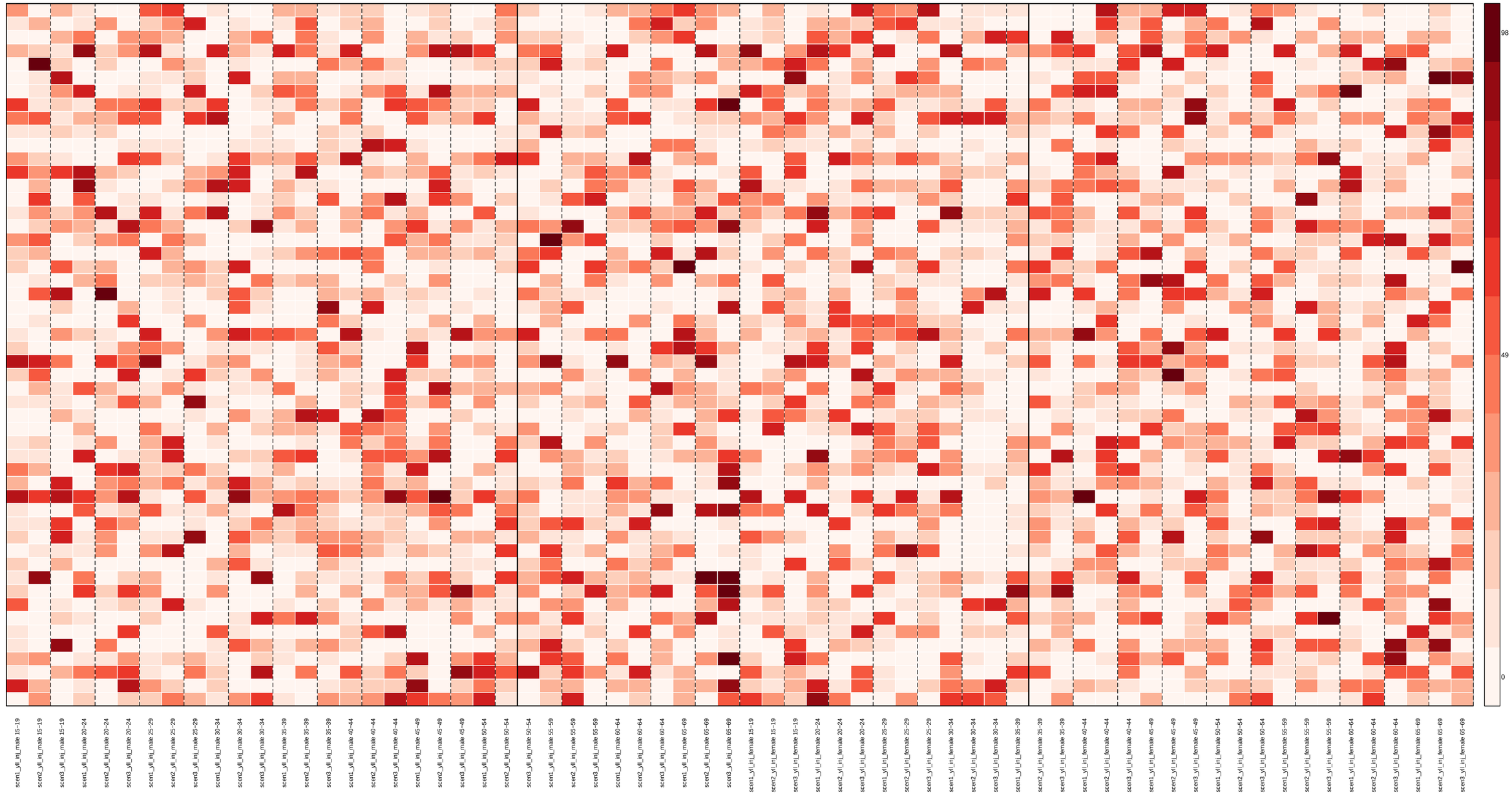
montevideo − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



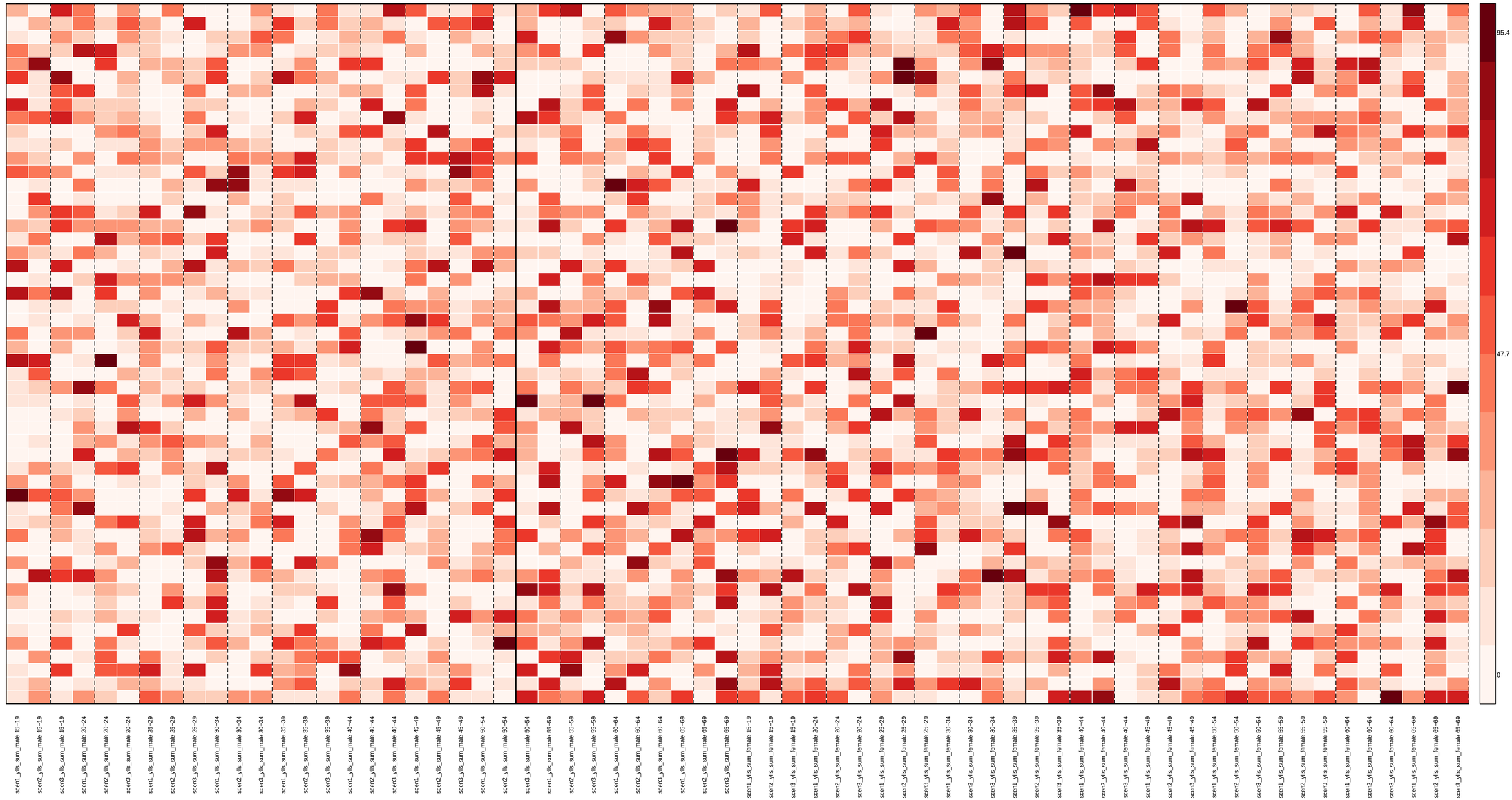
montevideo − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



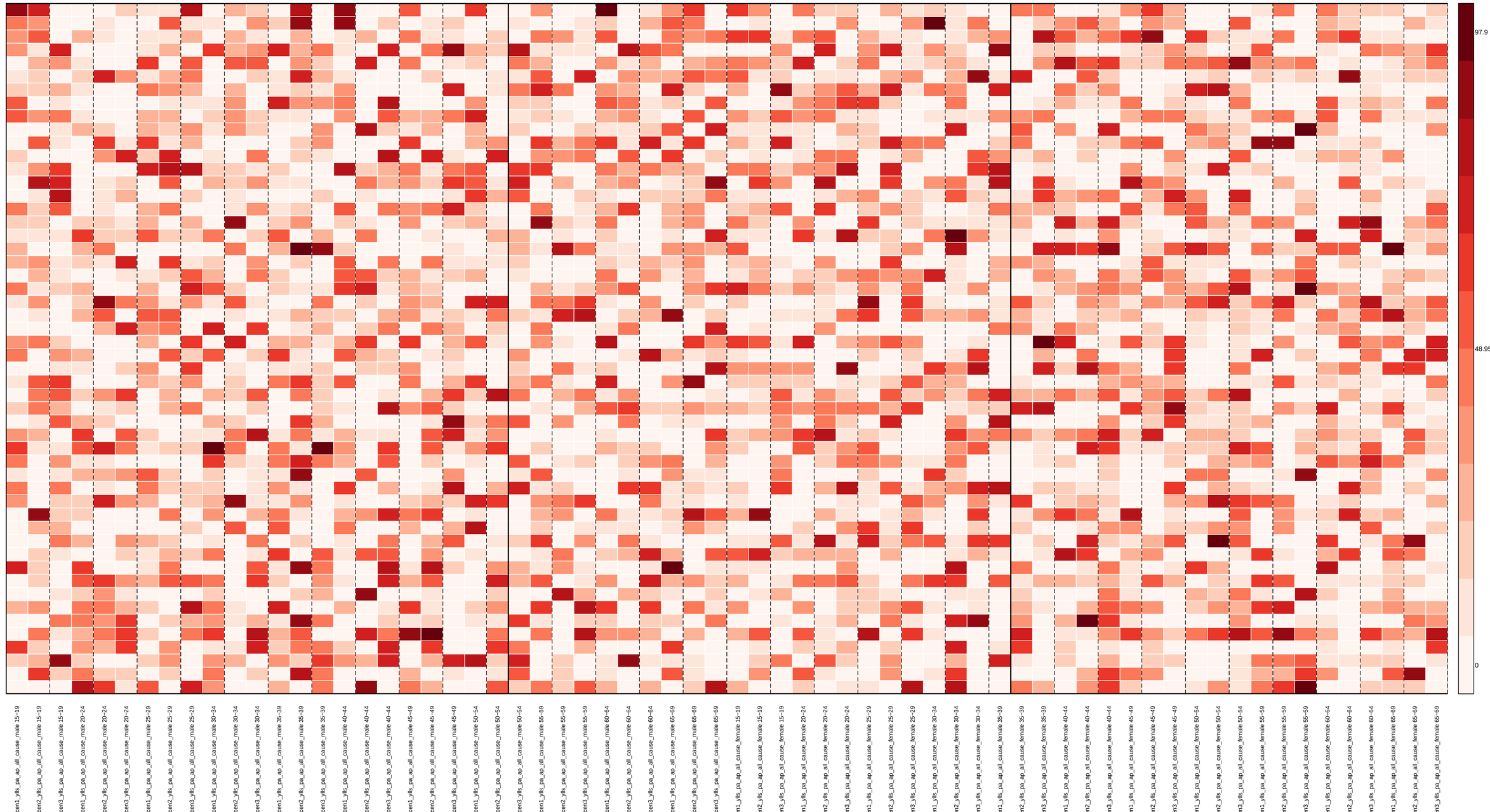
montevideo − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



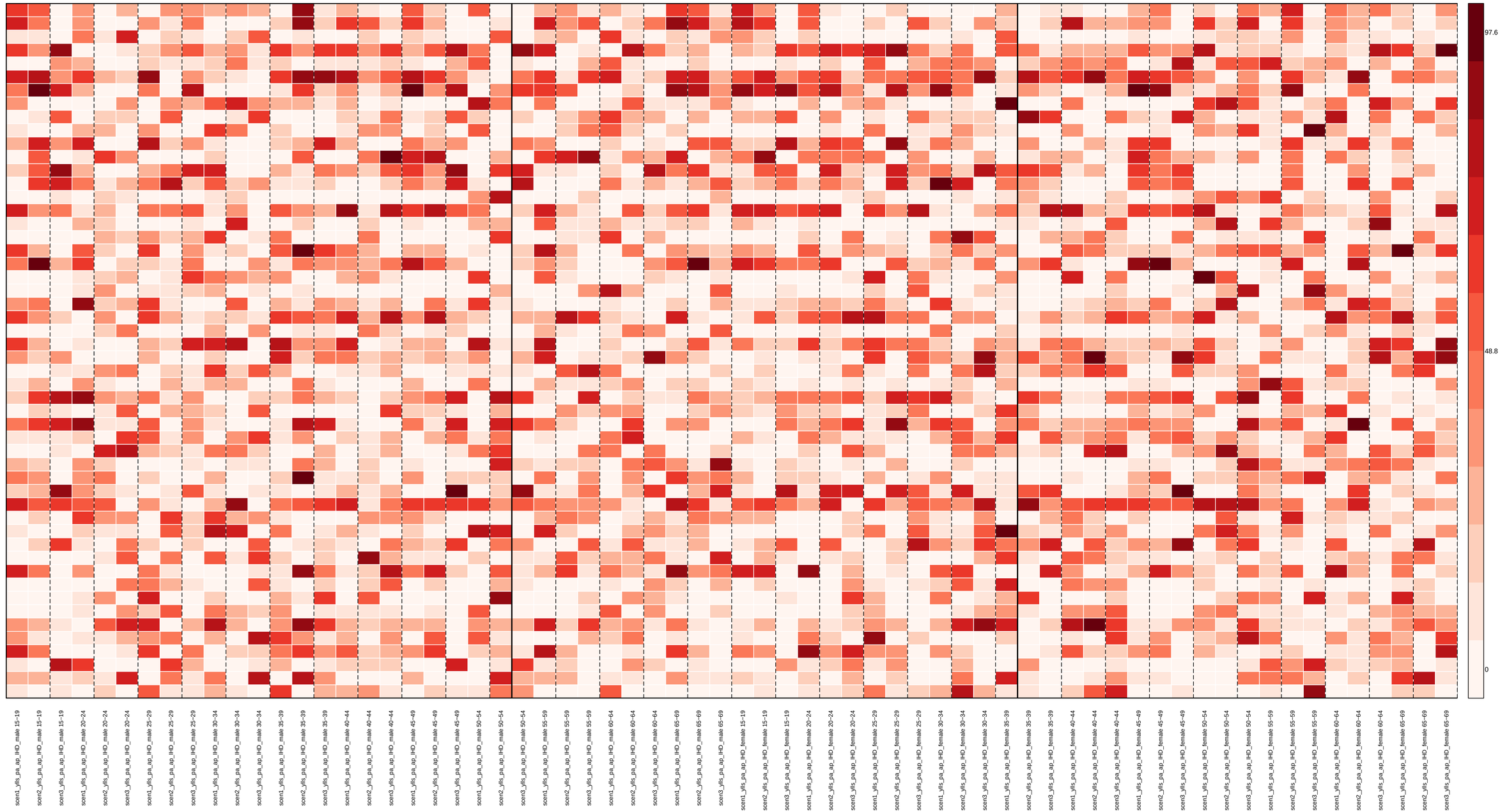
montevideo − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



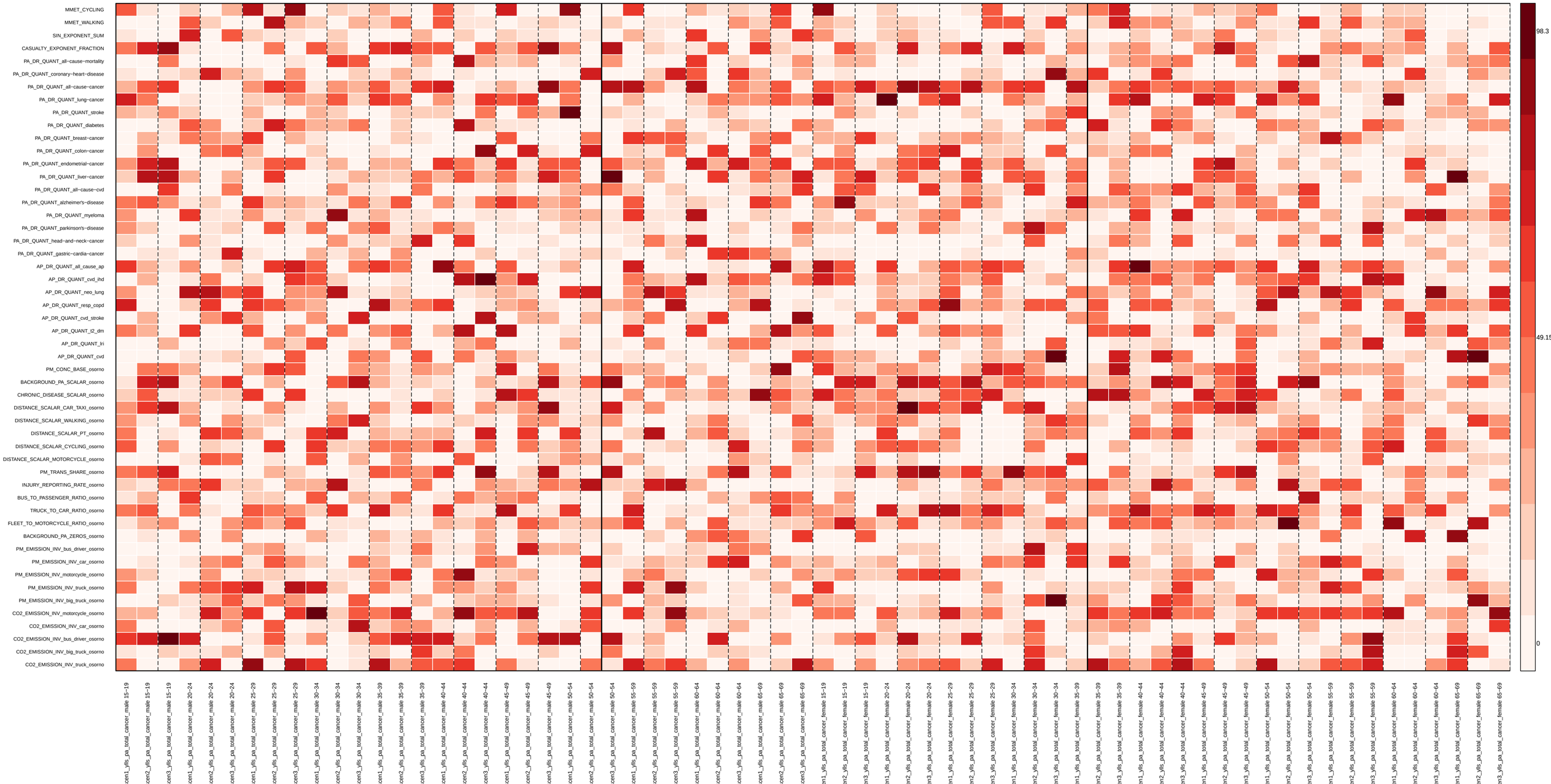
osorno − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



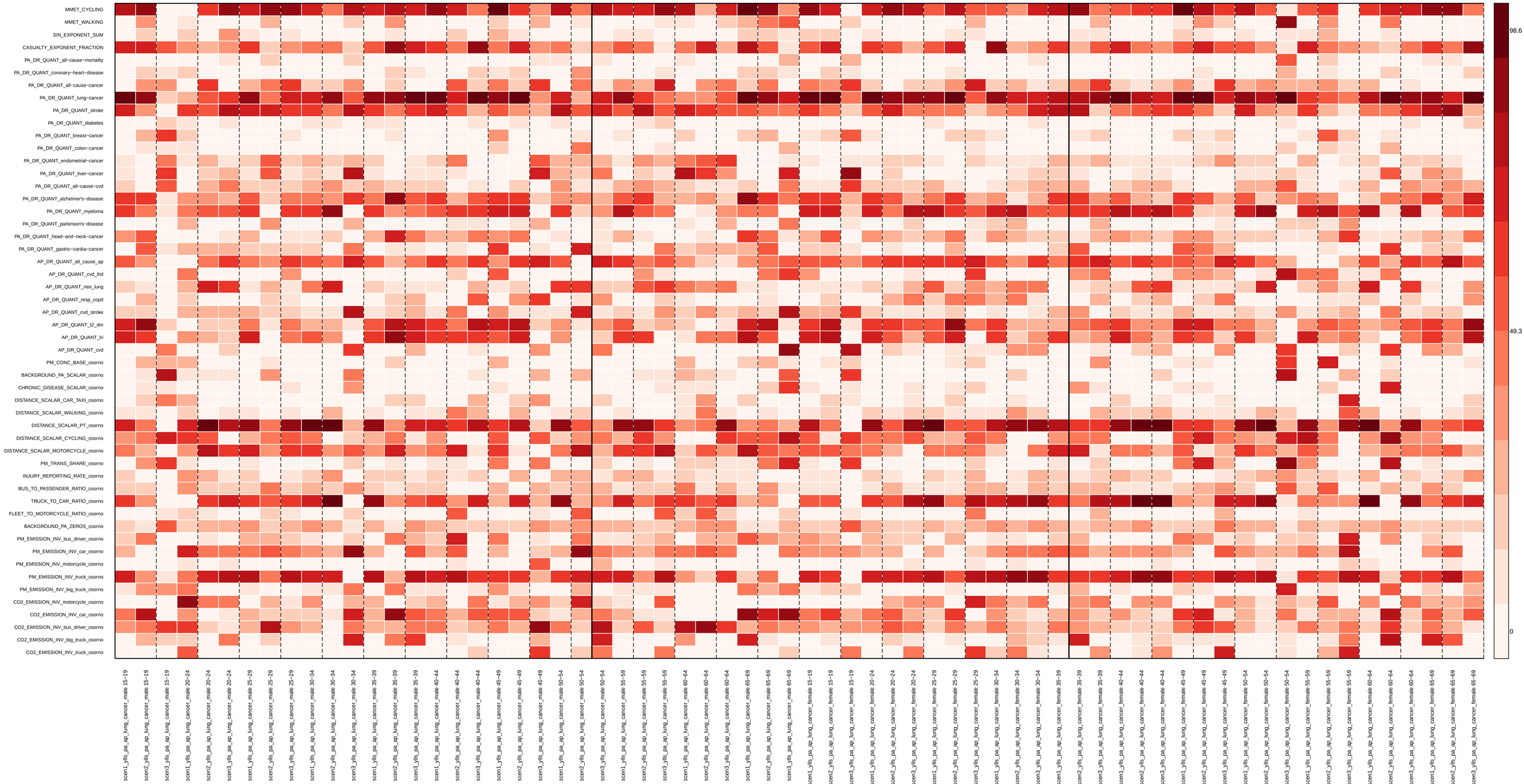
osorno − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



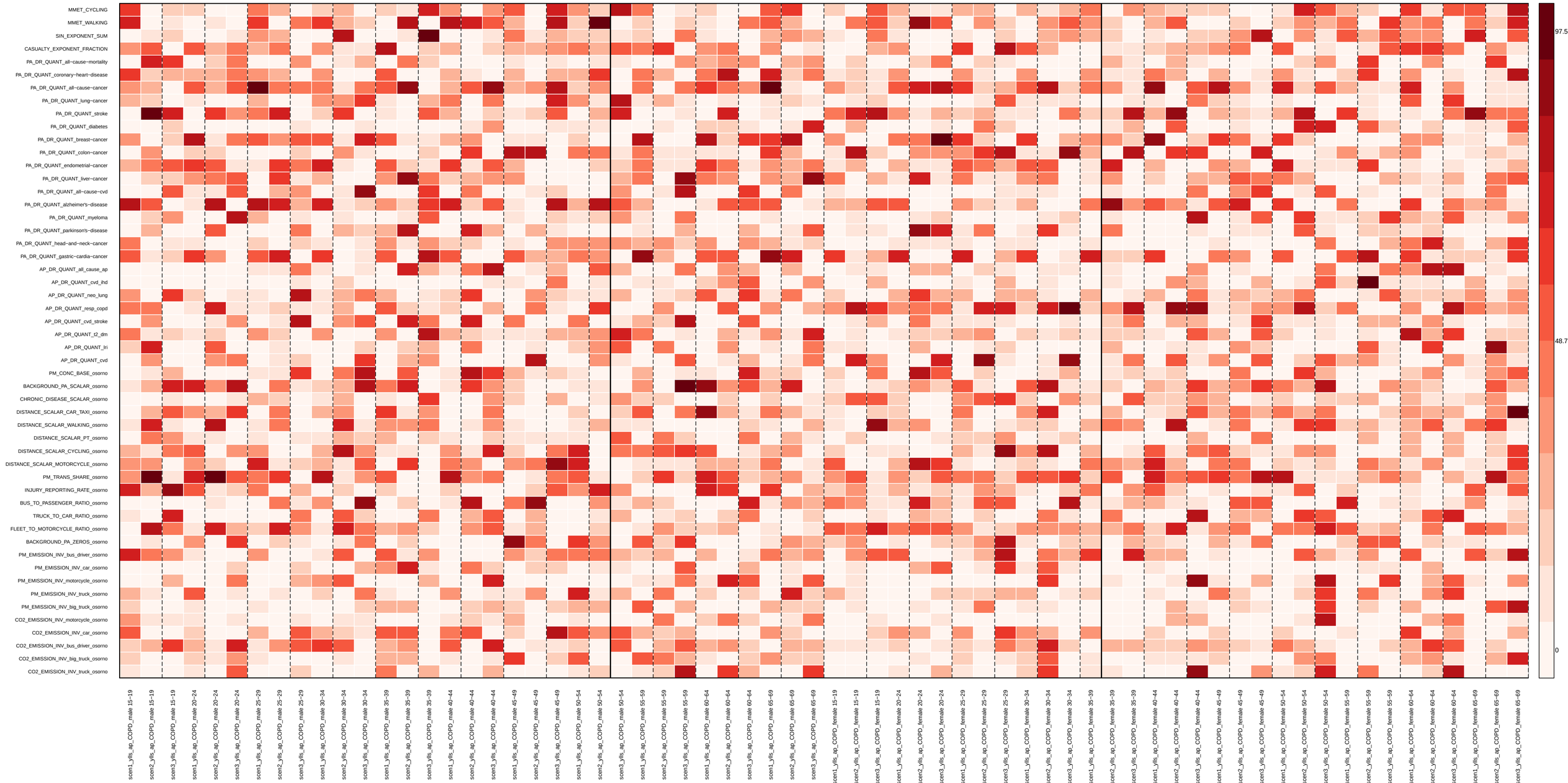
osorno − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



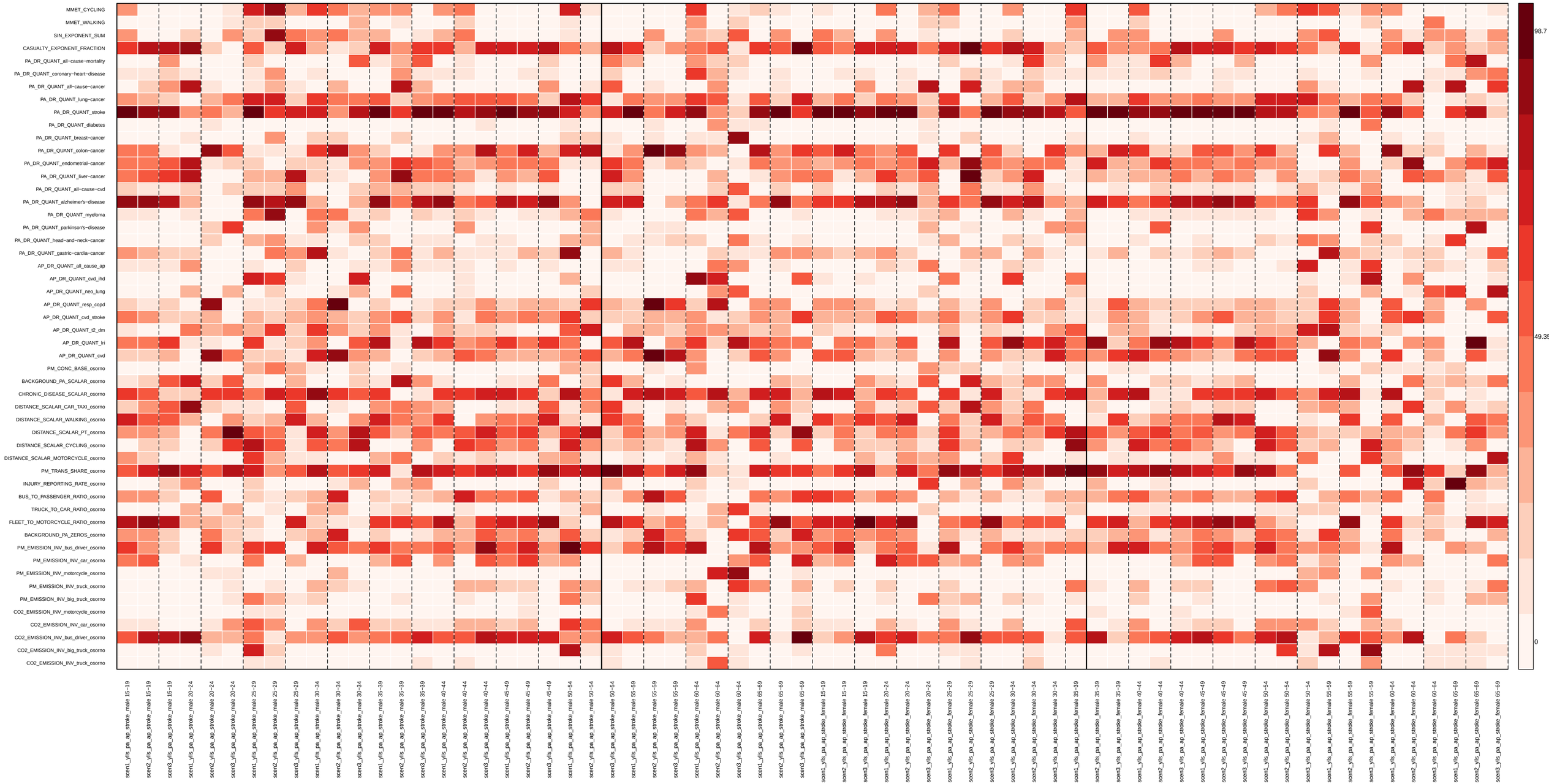
osorno − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



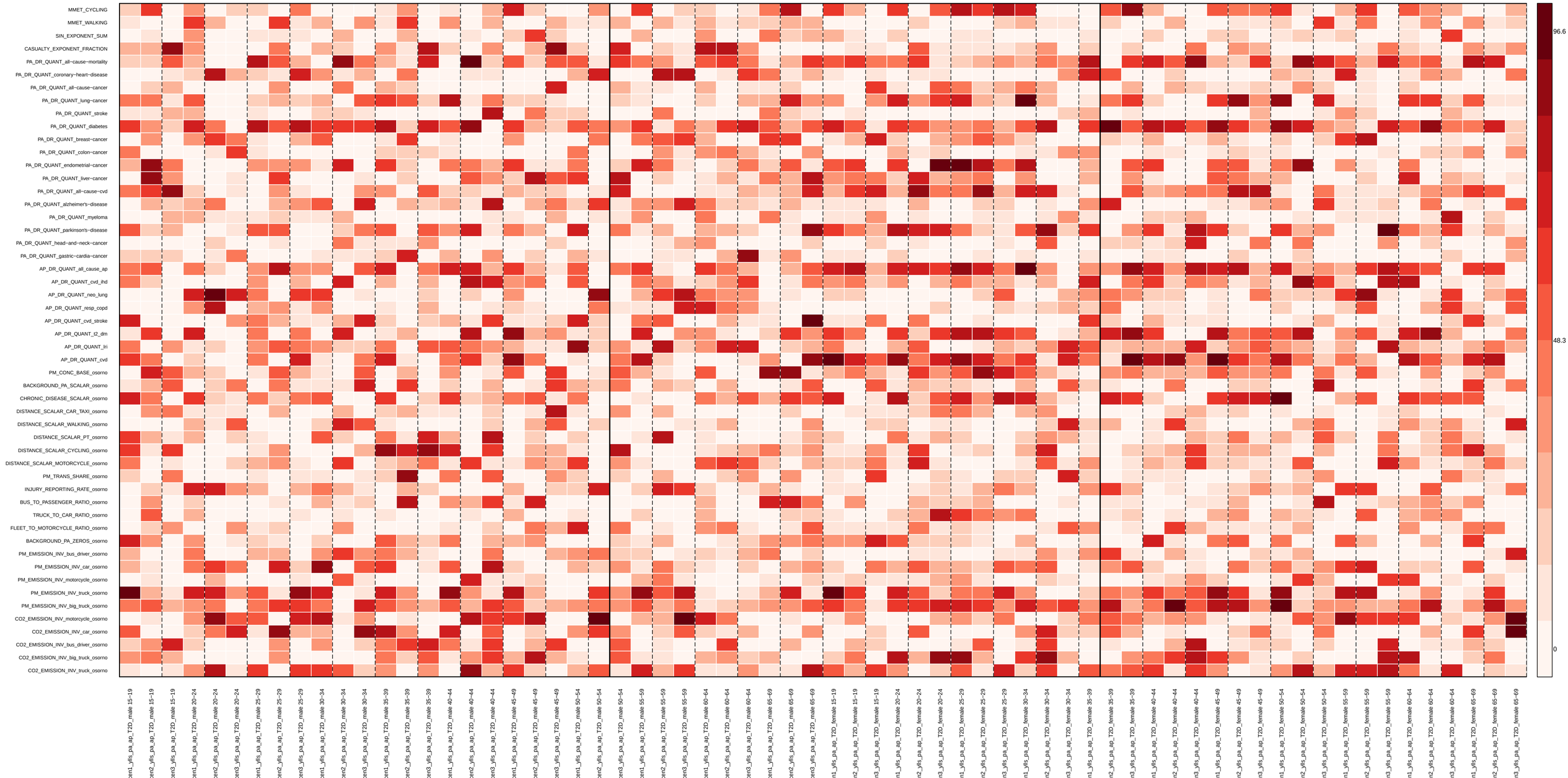
osorno − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



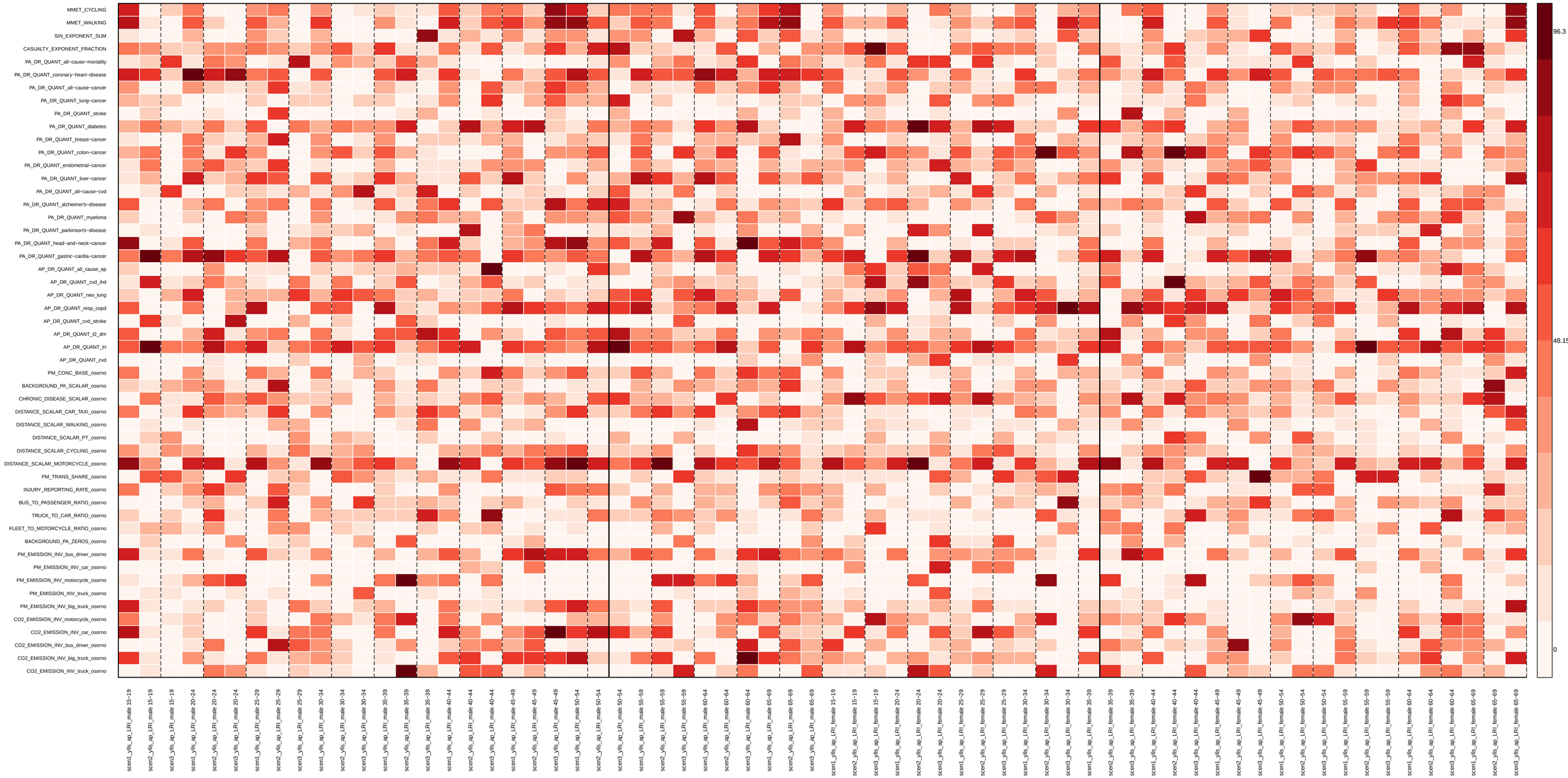
osorno − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



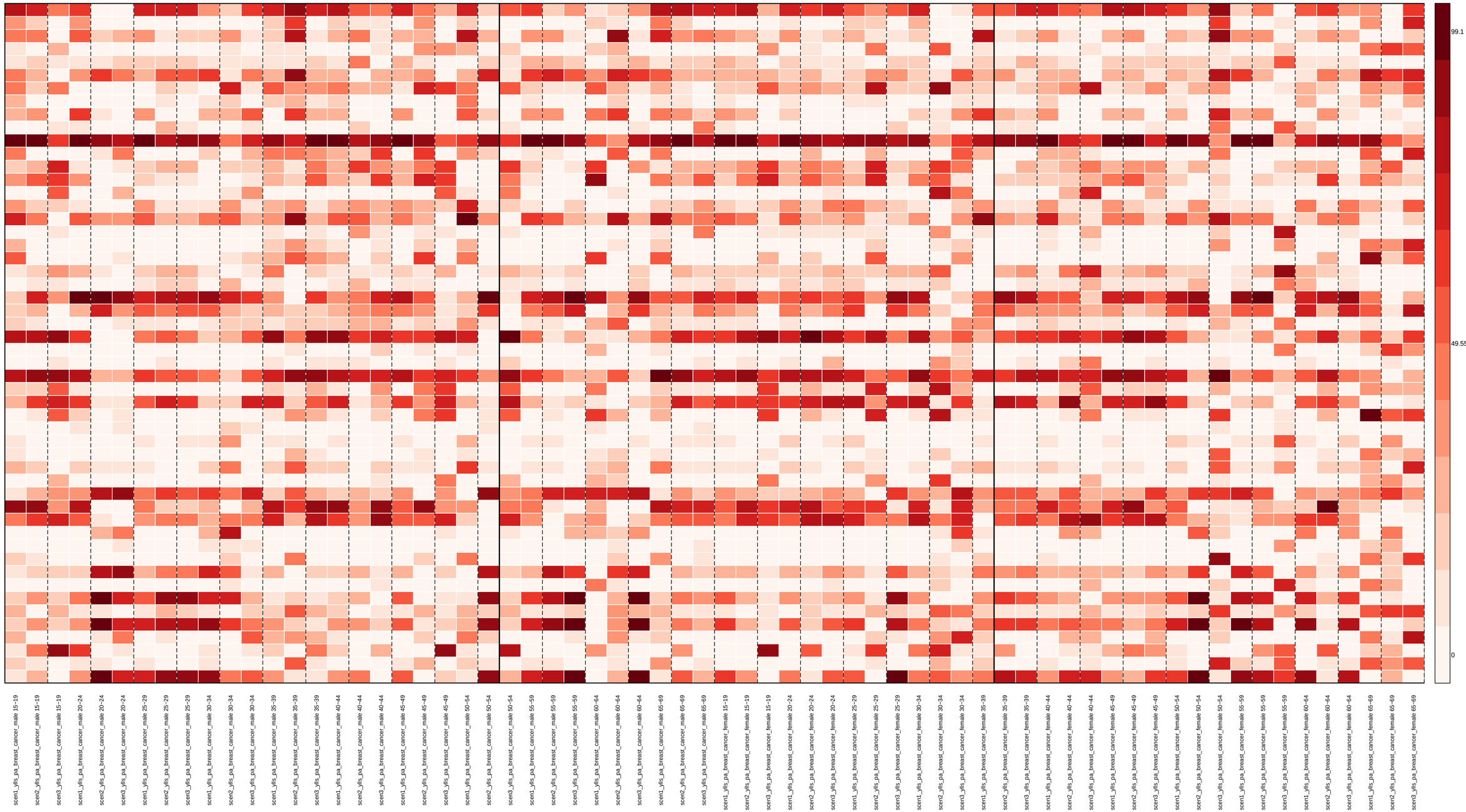
osorno − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



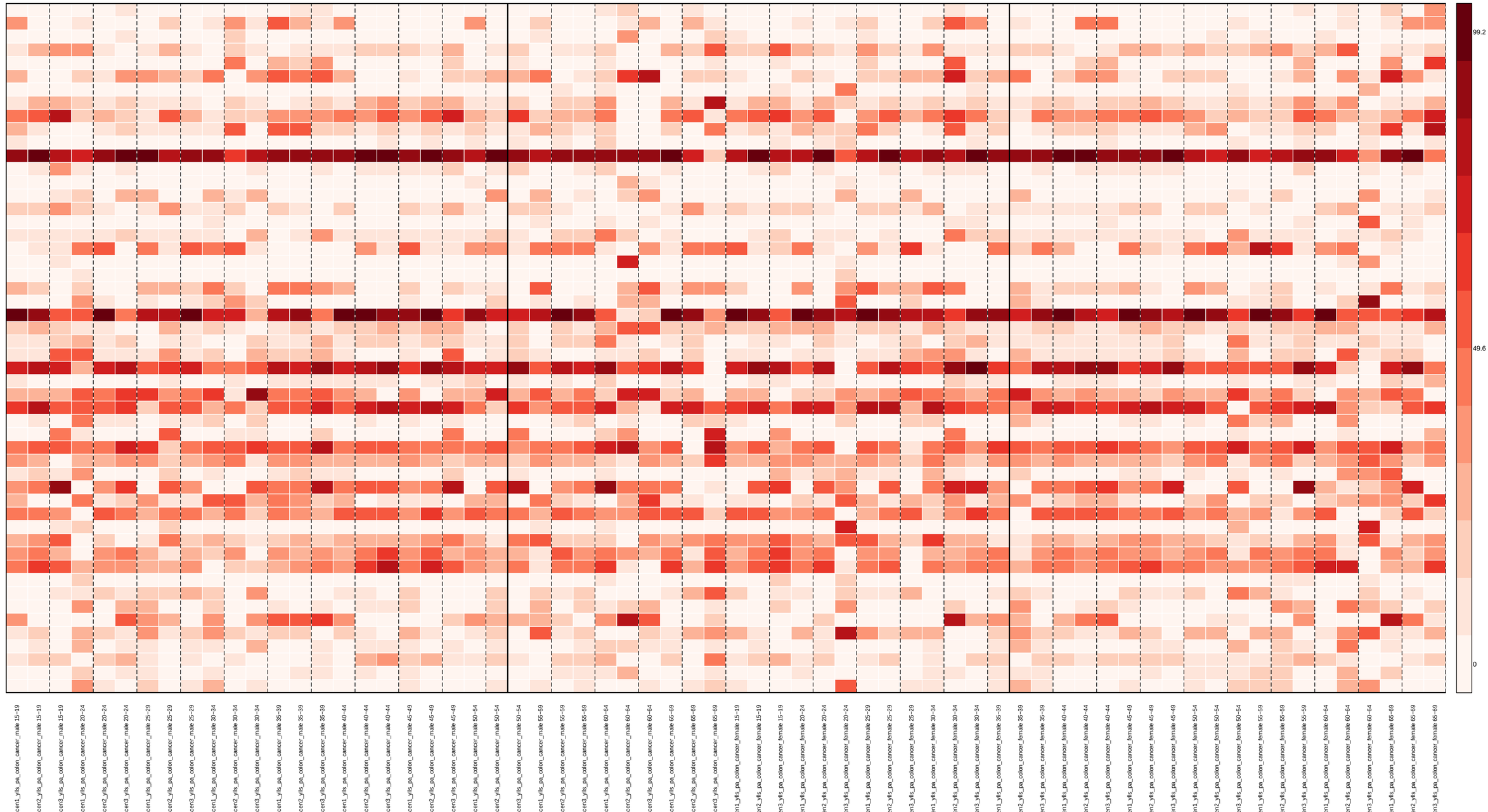
osorno − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



osorno − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

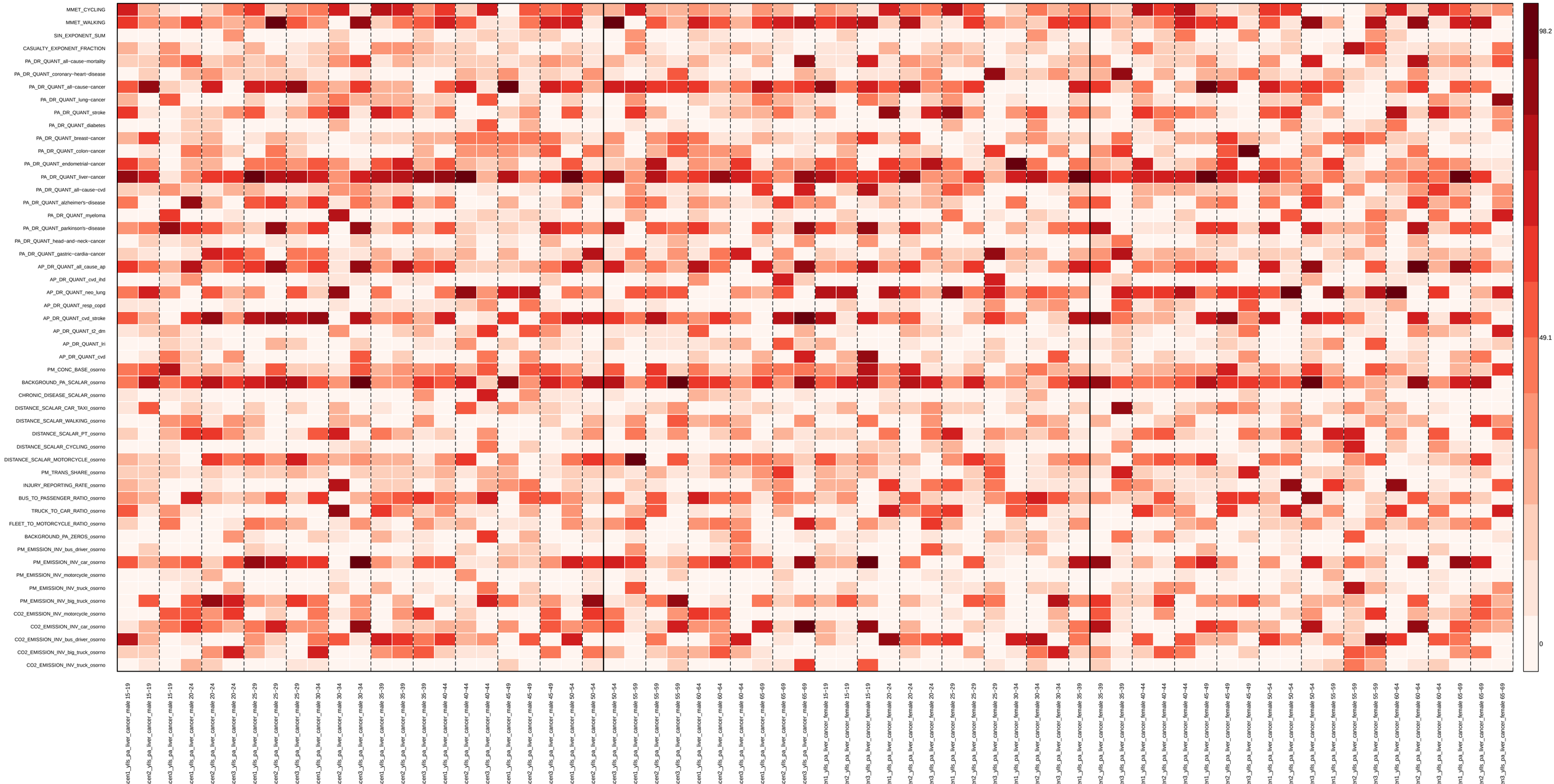


osorno − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

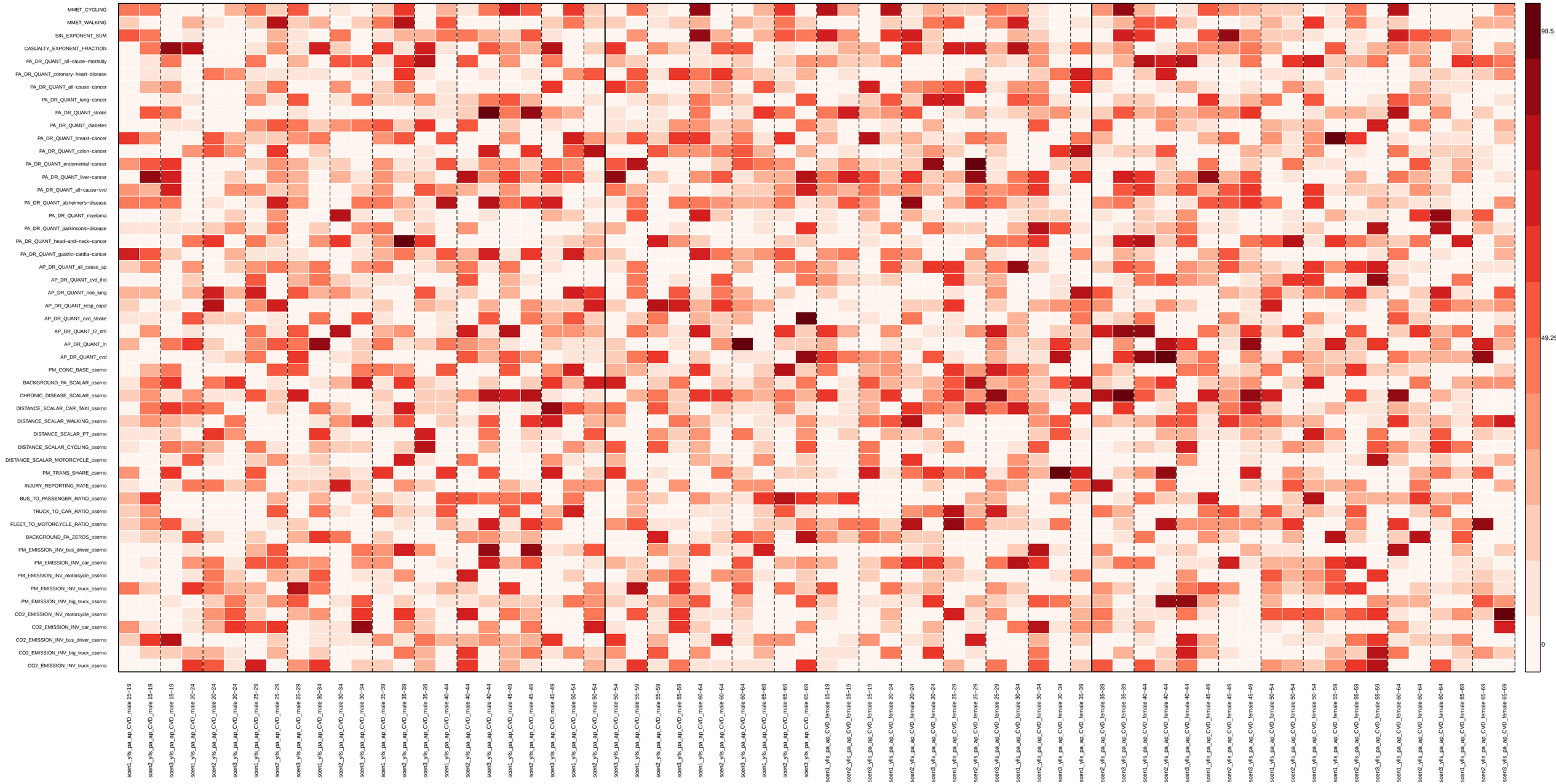


osorno − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

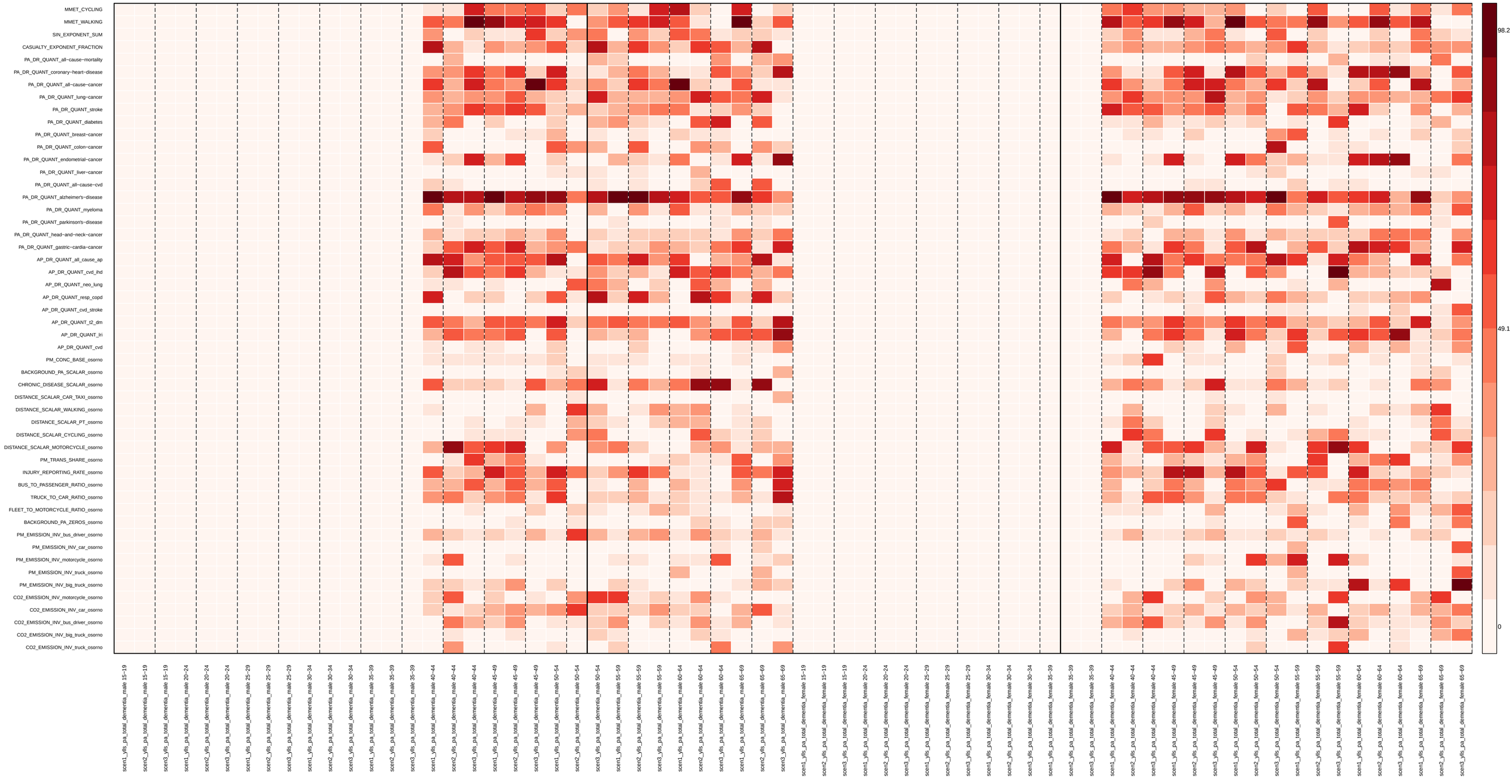
osorno − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



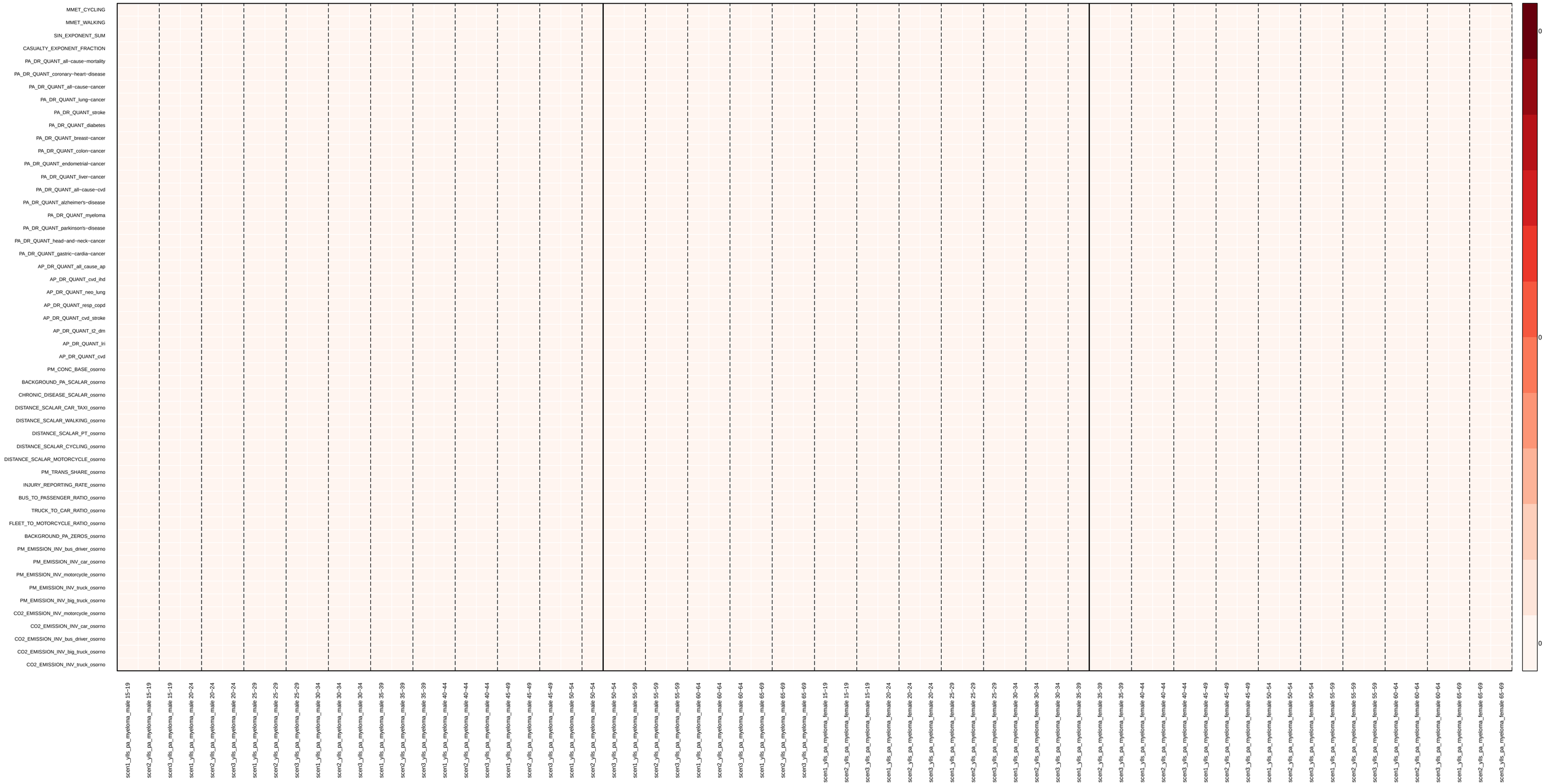
osorno − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



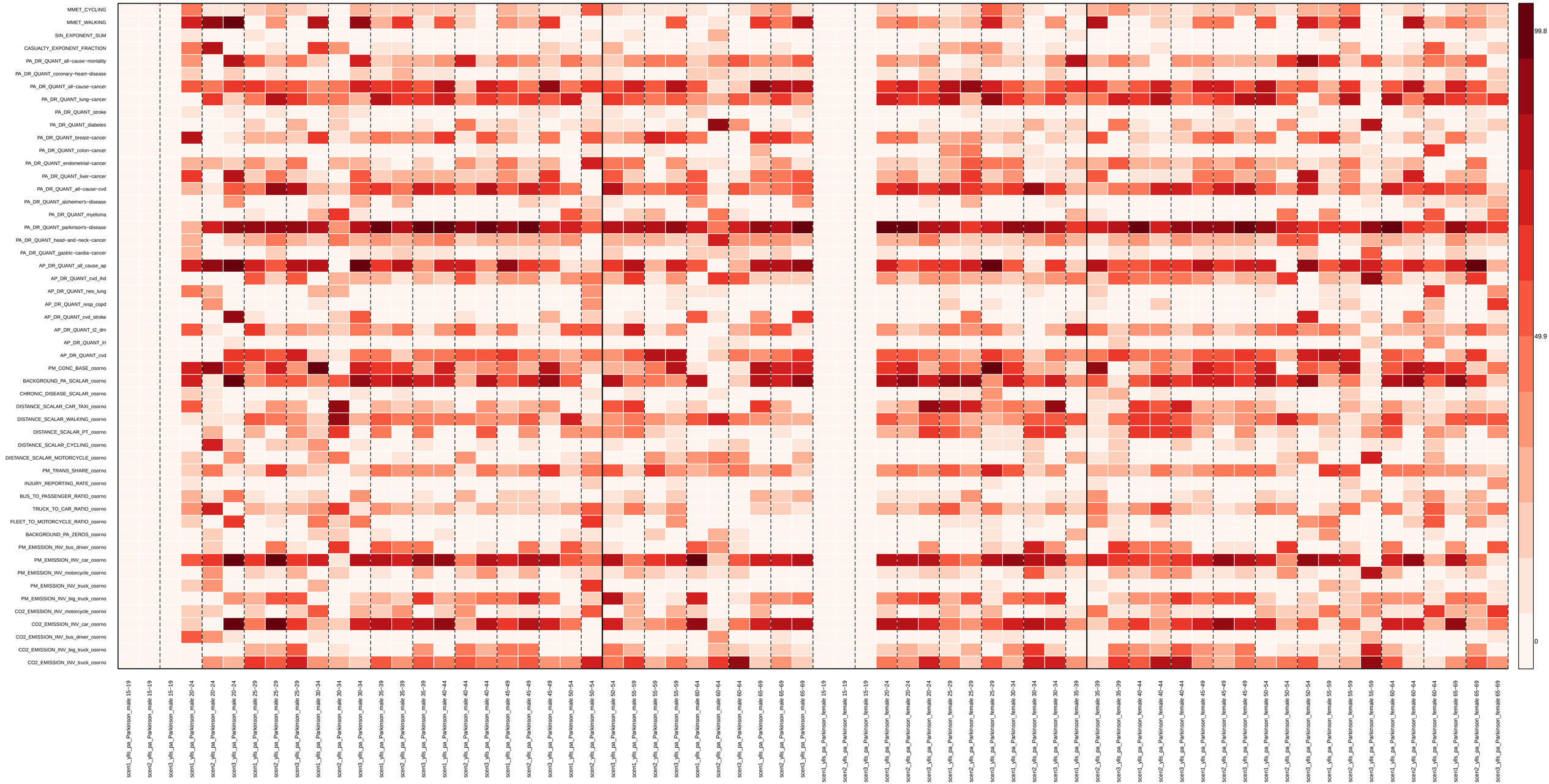
osorno − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



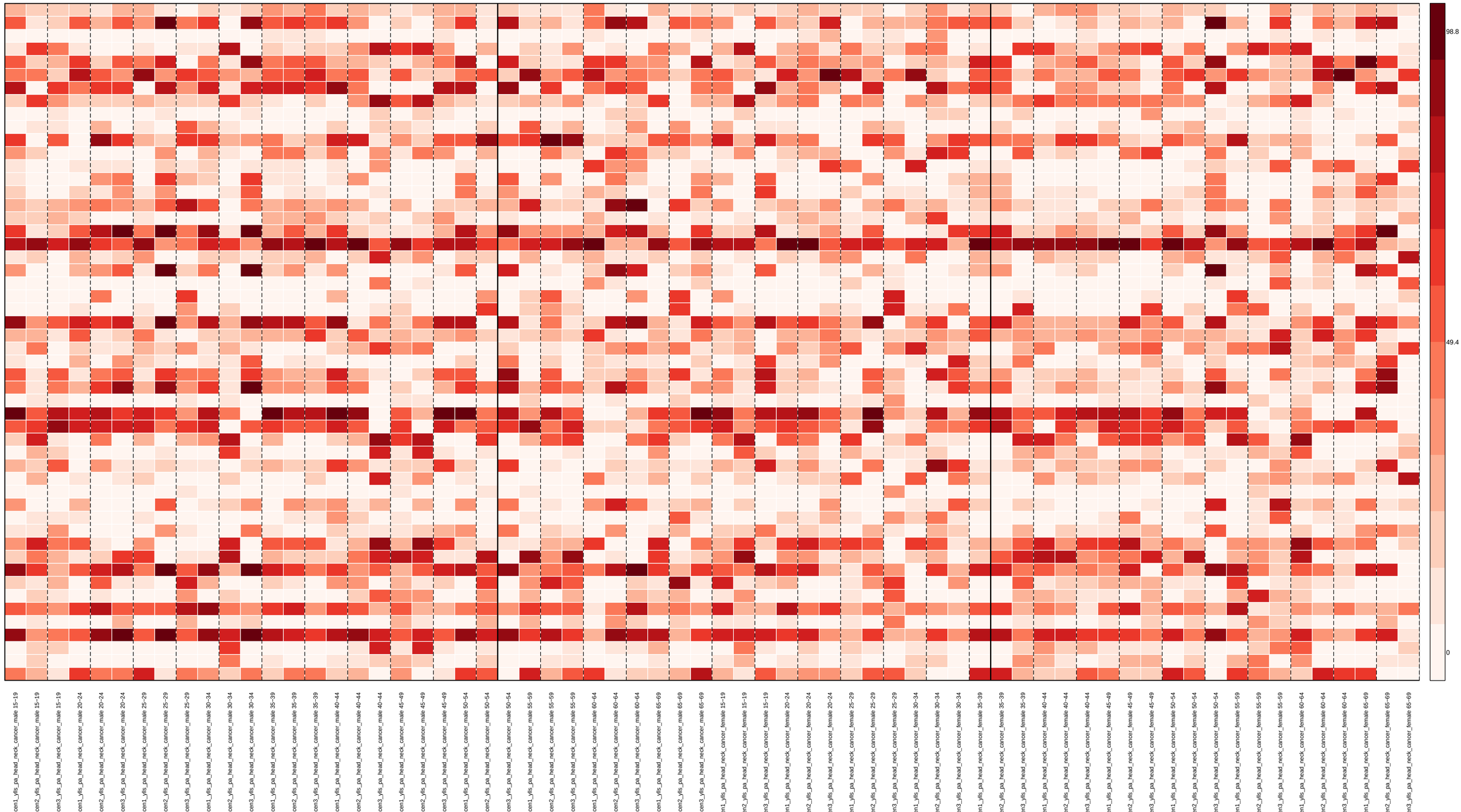
osorno − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



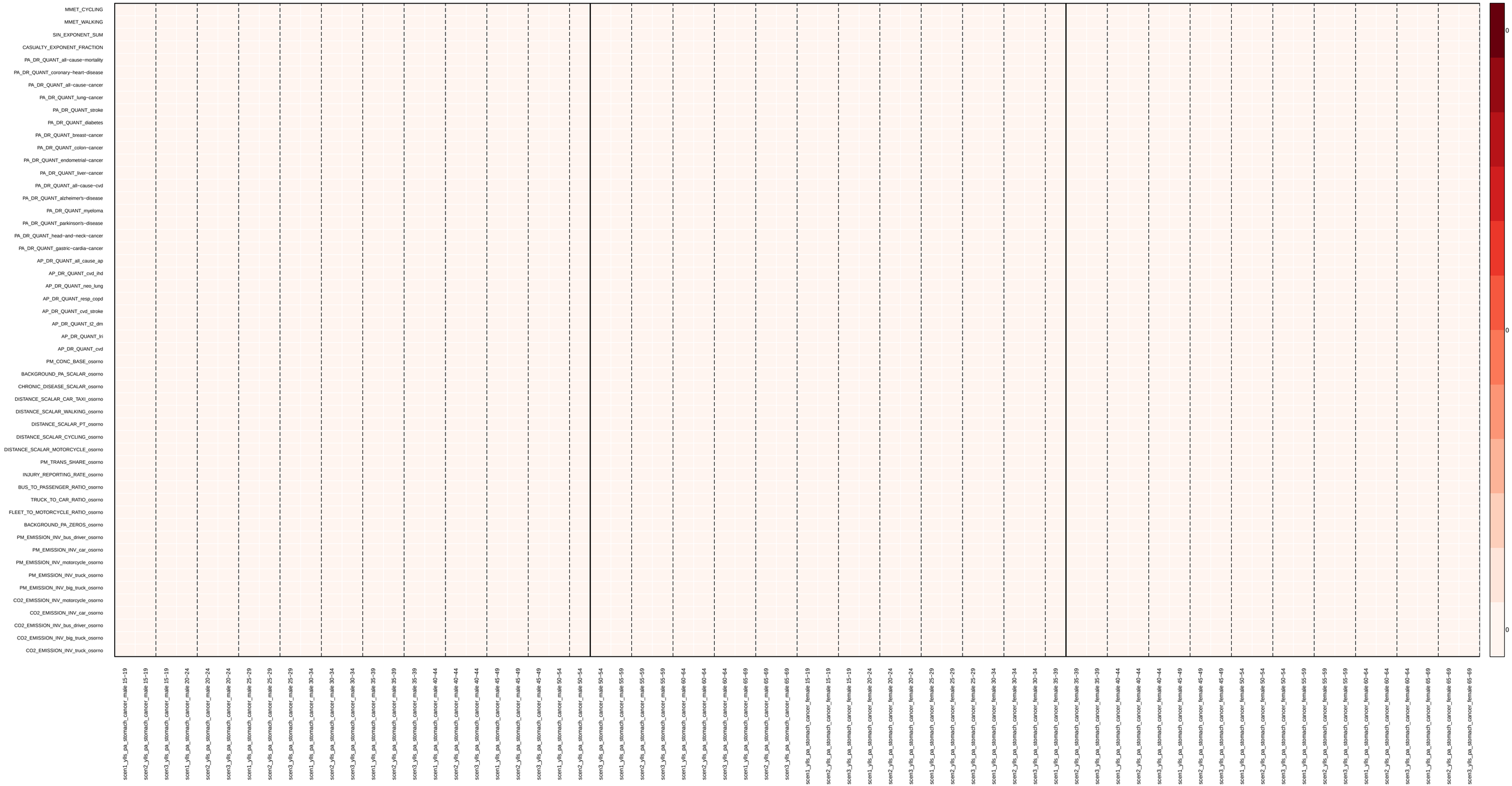
osorno − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



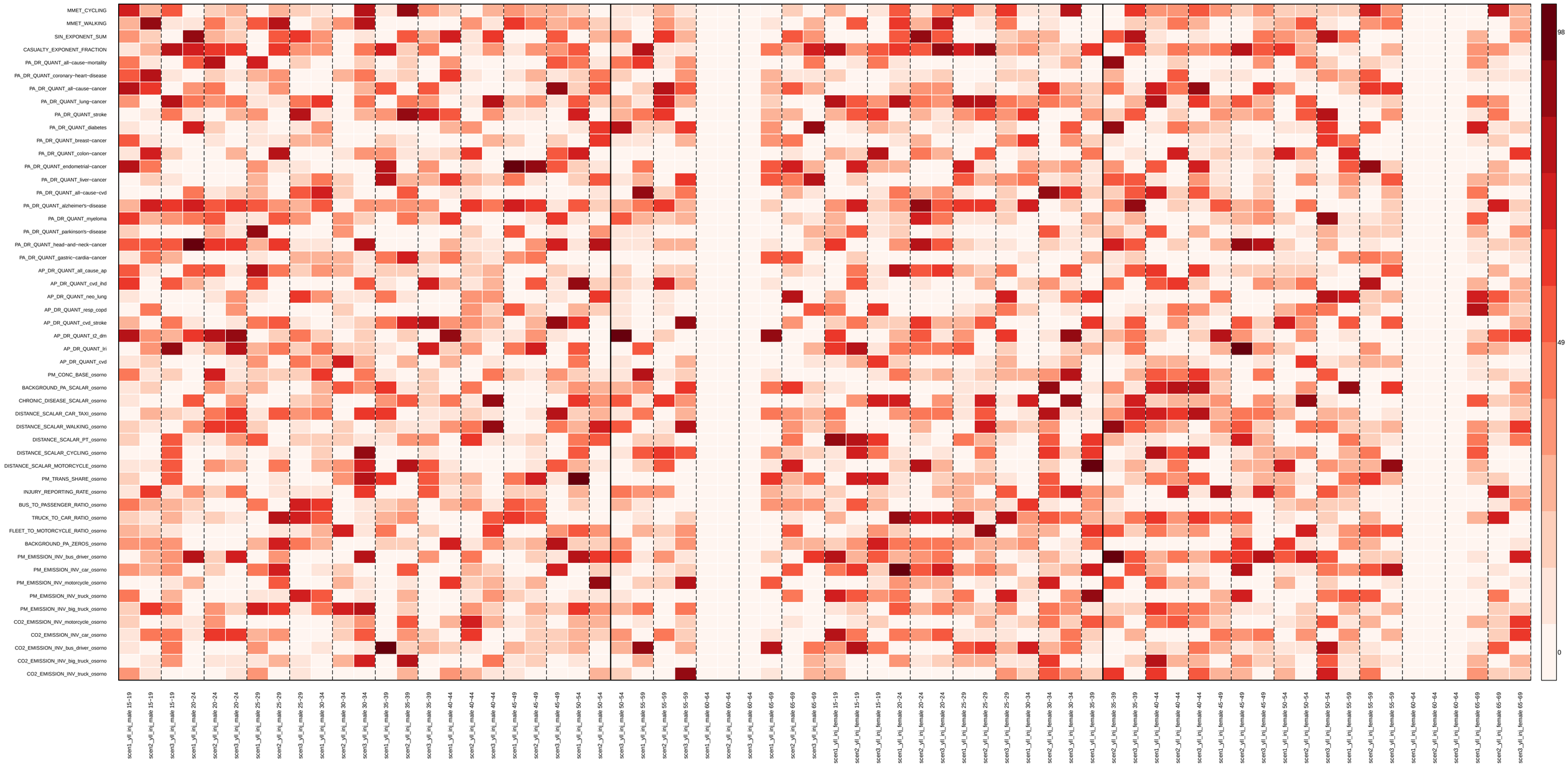
osorno − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



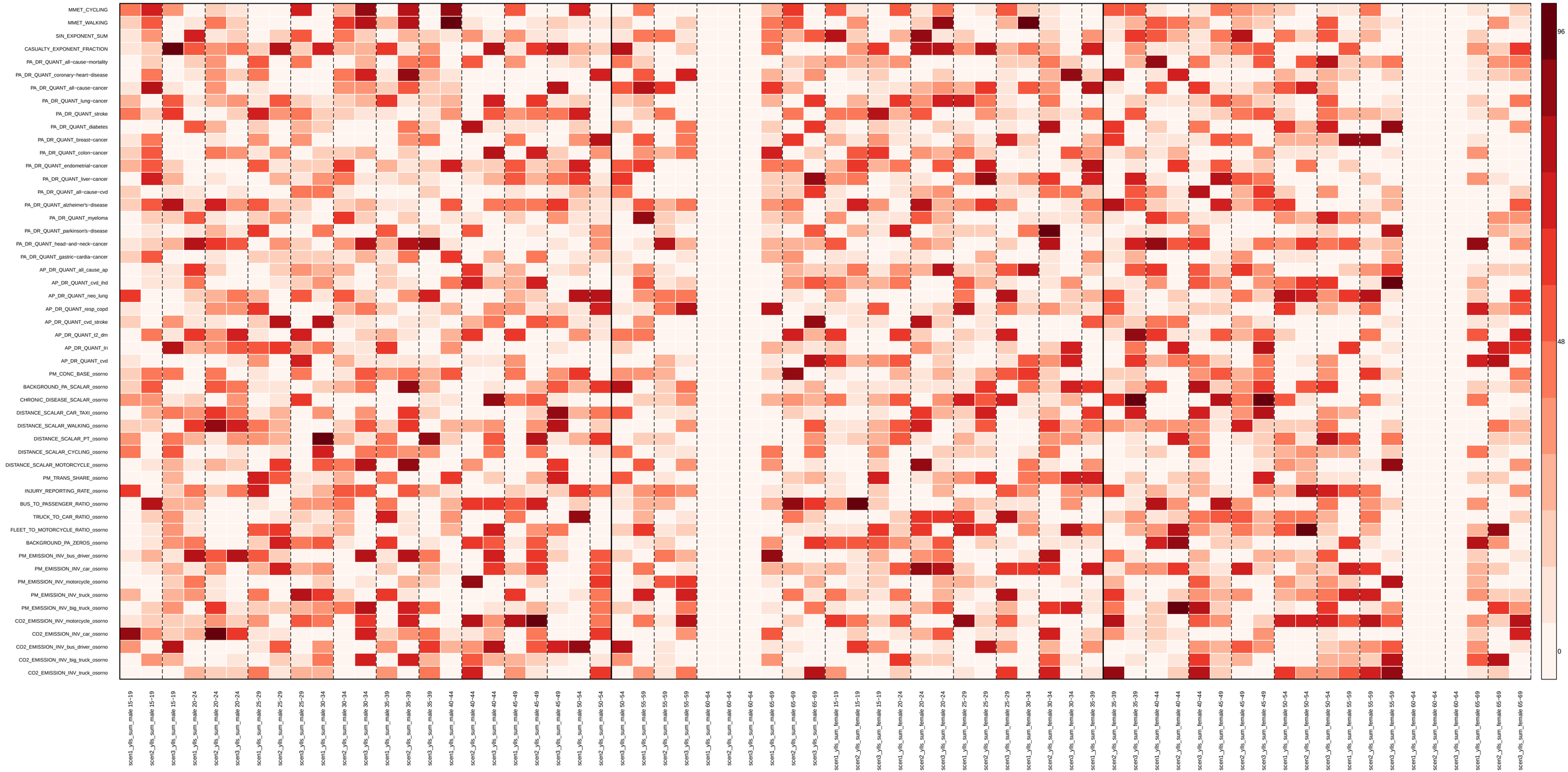
osorno − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



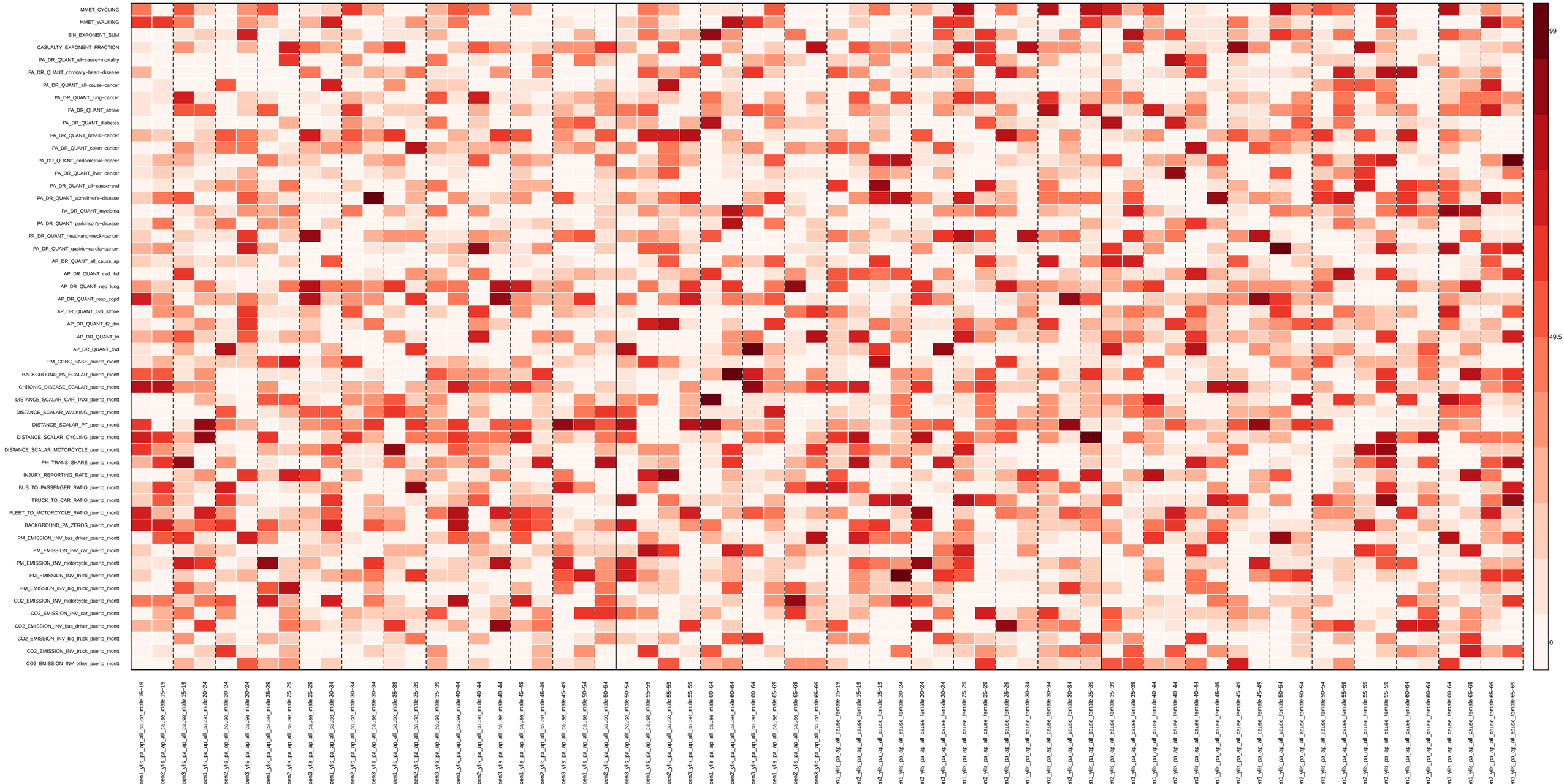
osorno − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



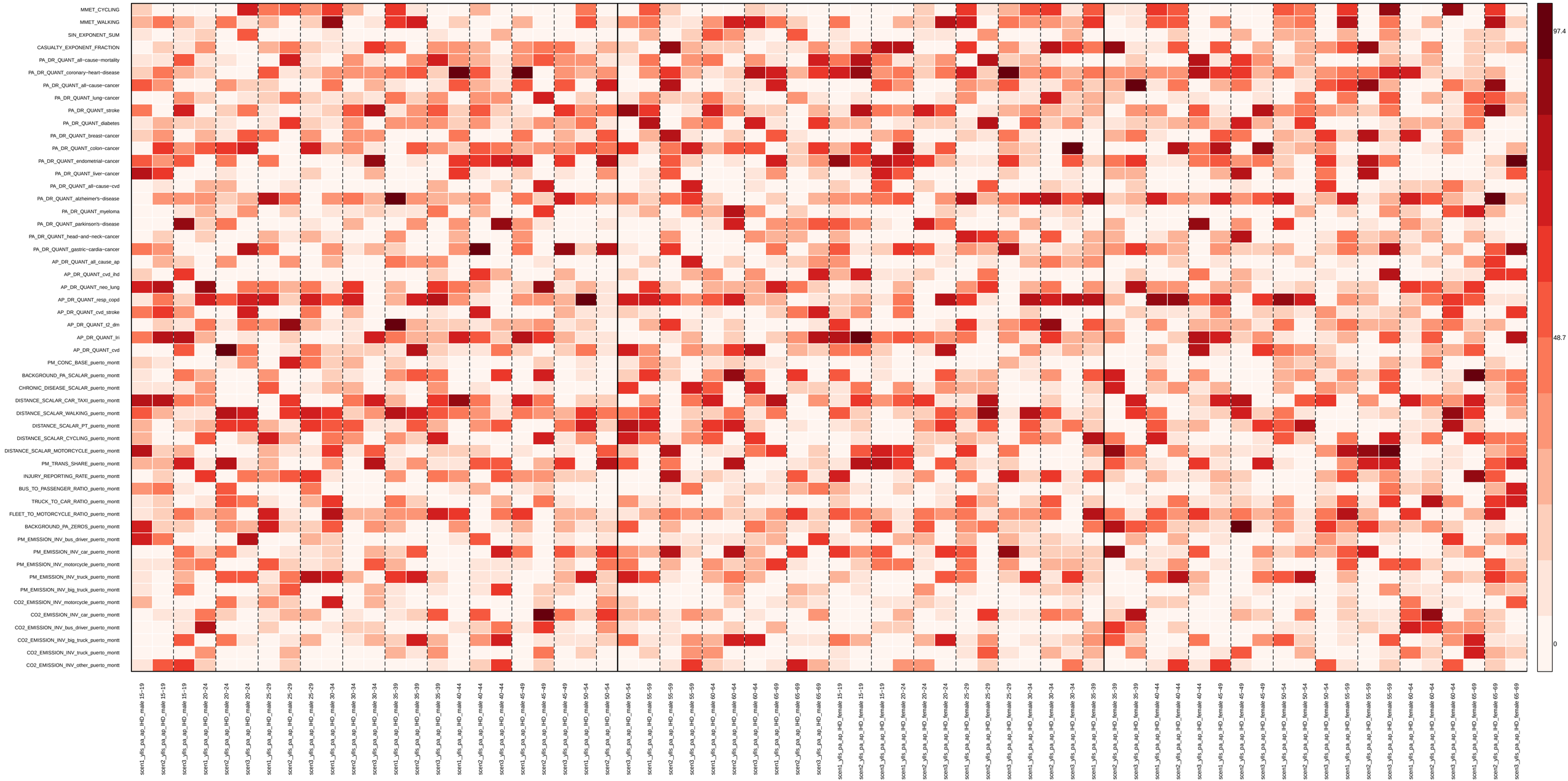
osorno − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



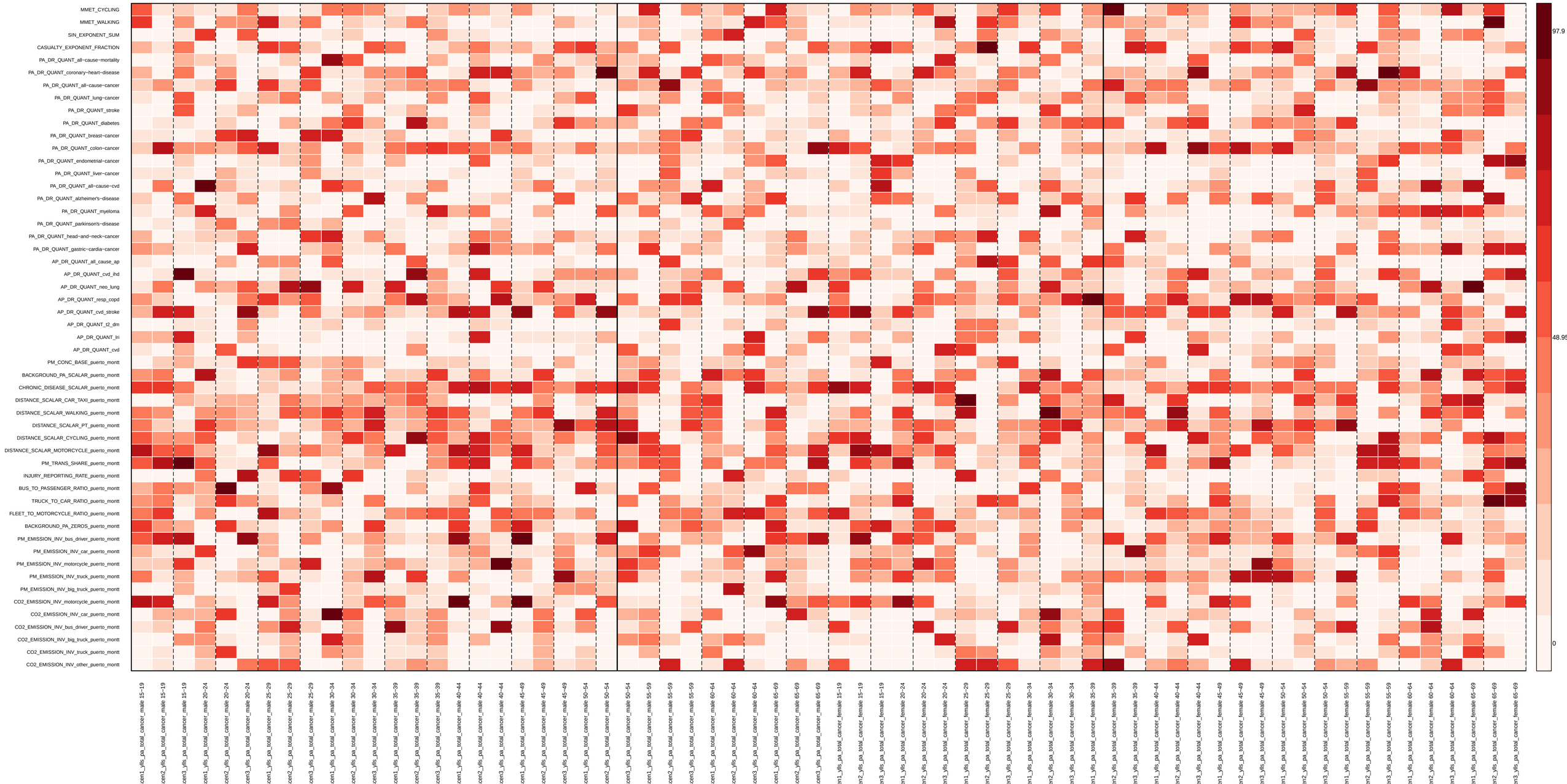
puerto_montt − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



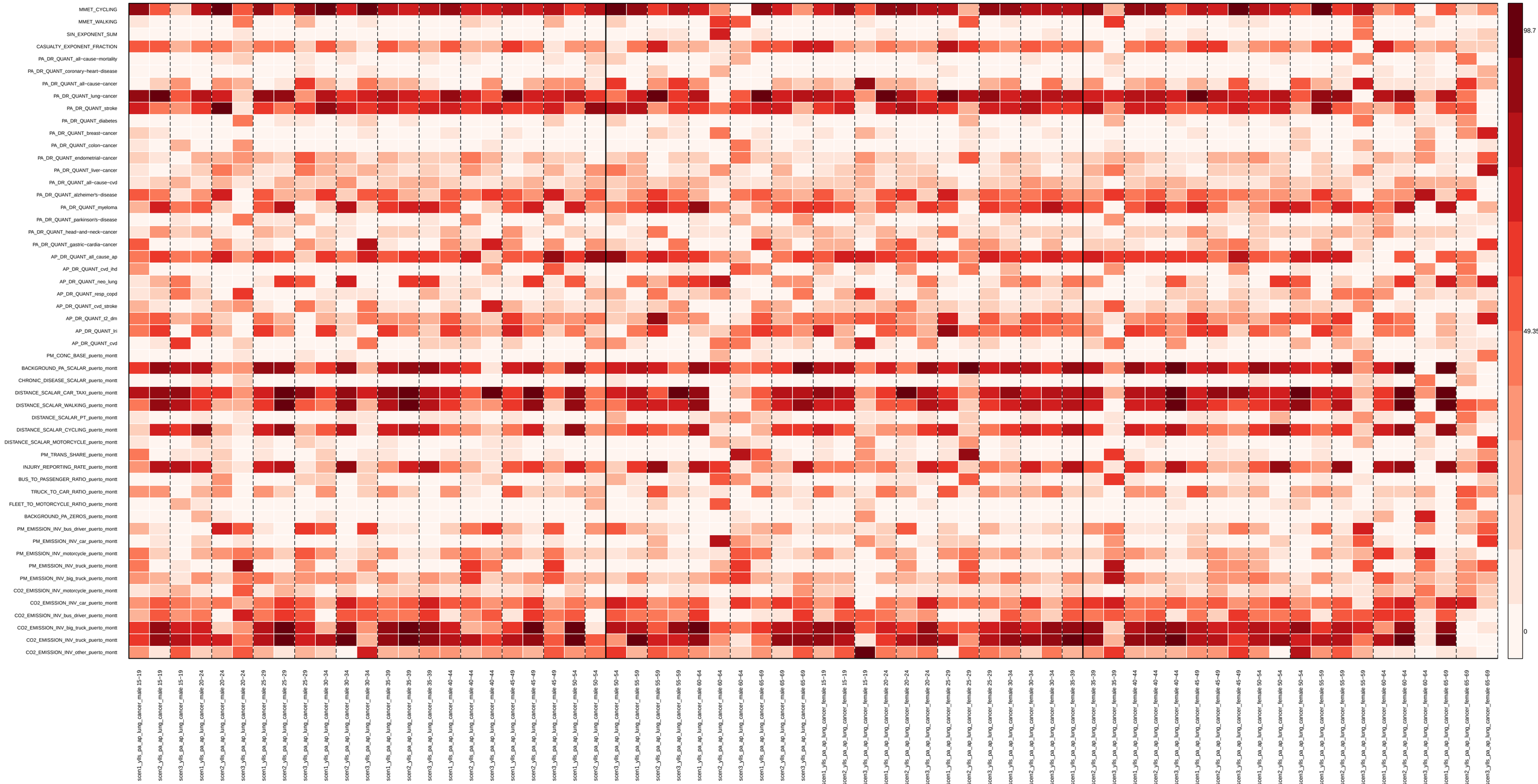
puerto_montt − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



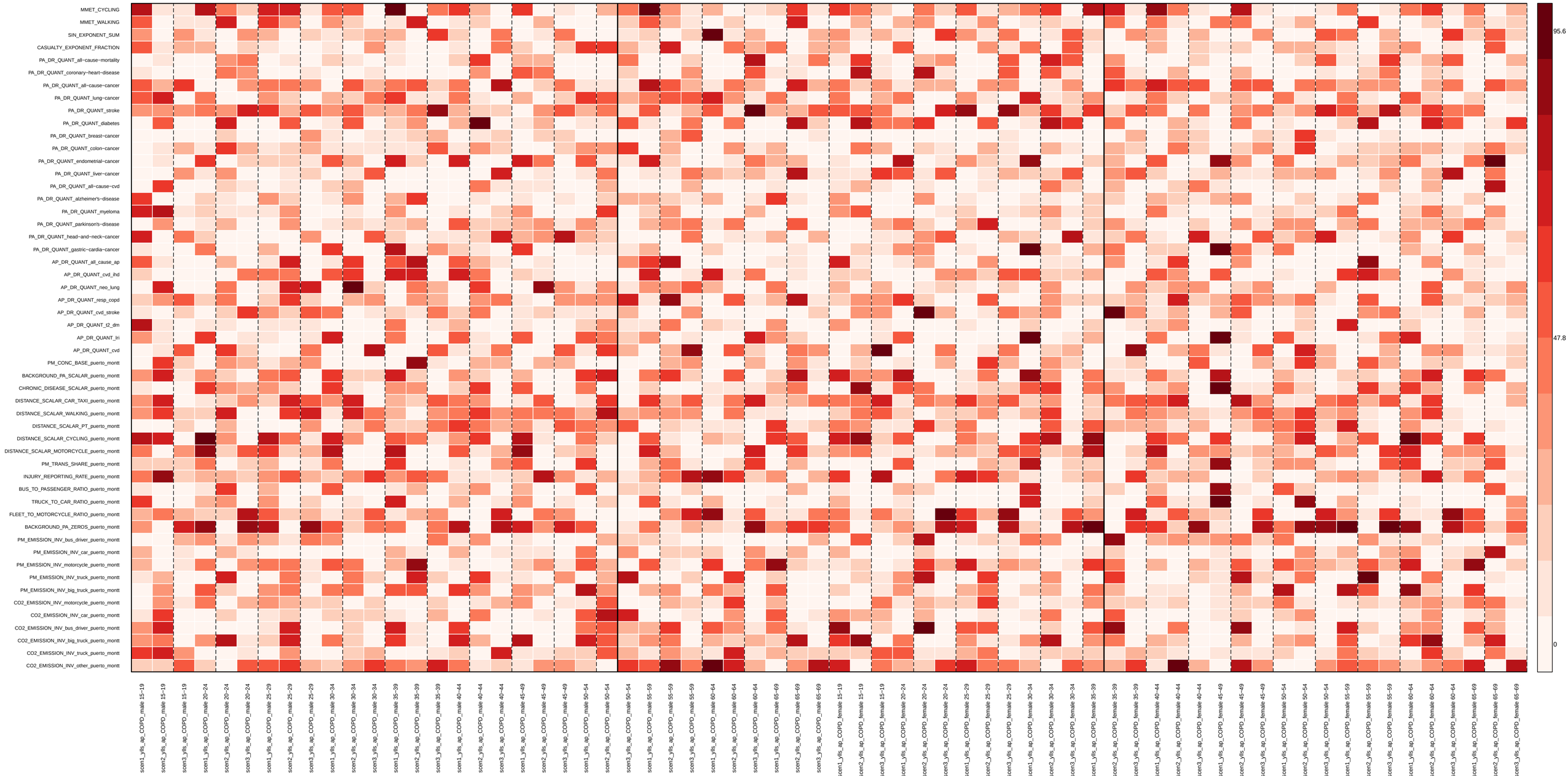
puerto_montt − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



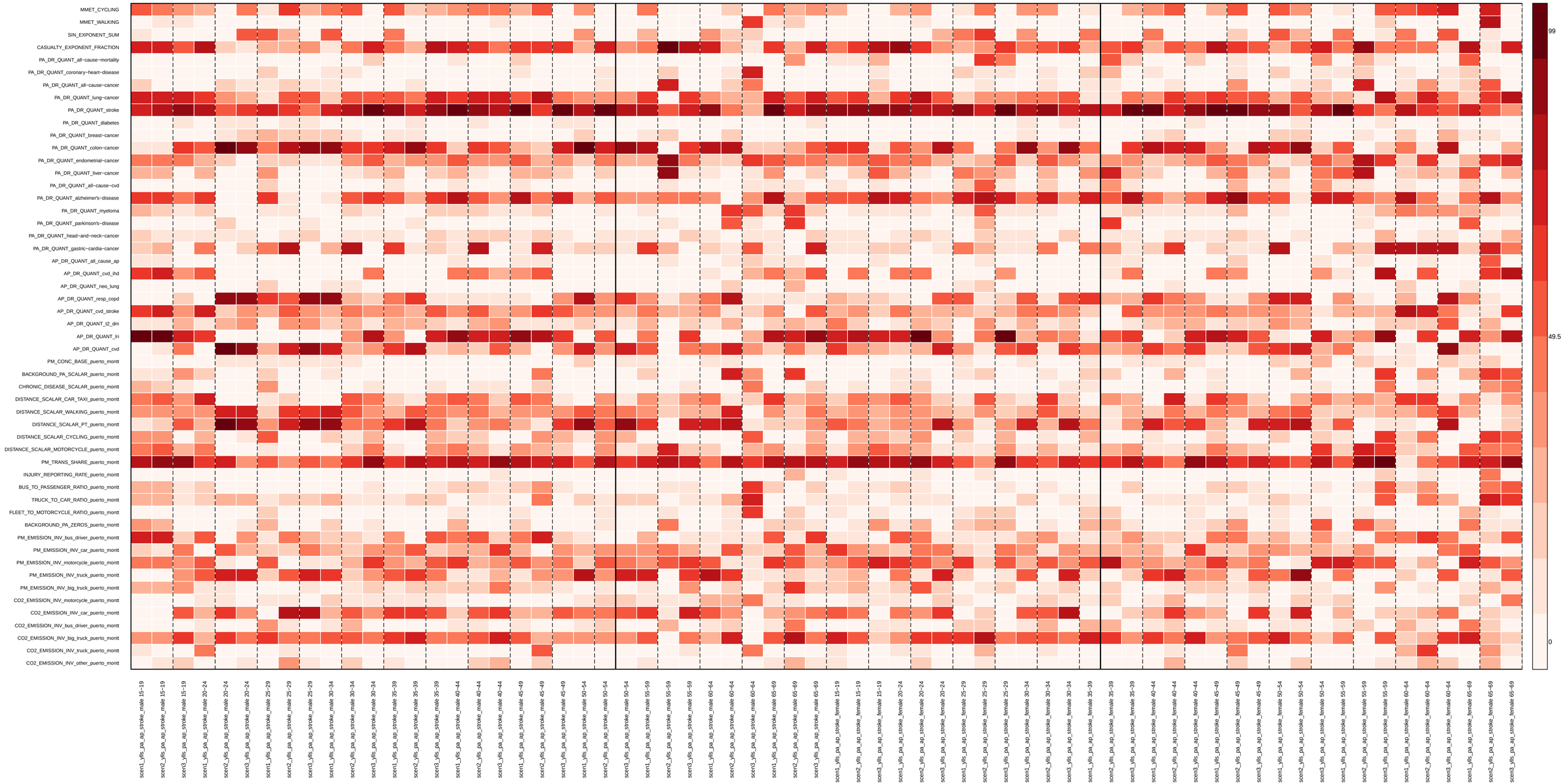
puerto_montt − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



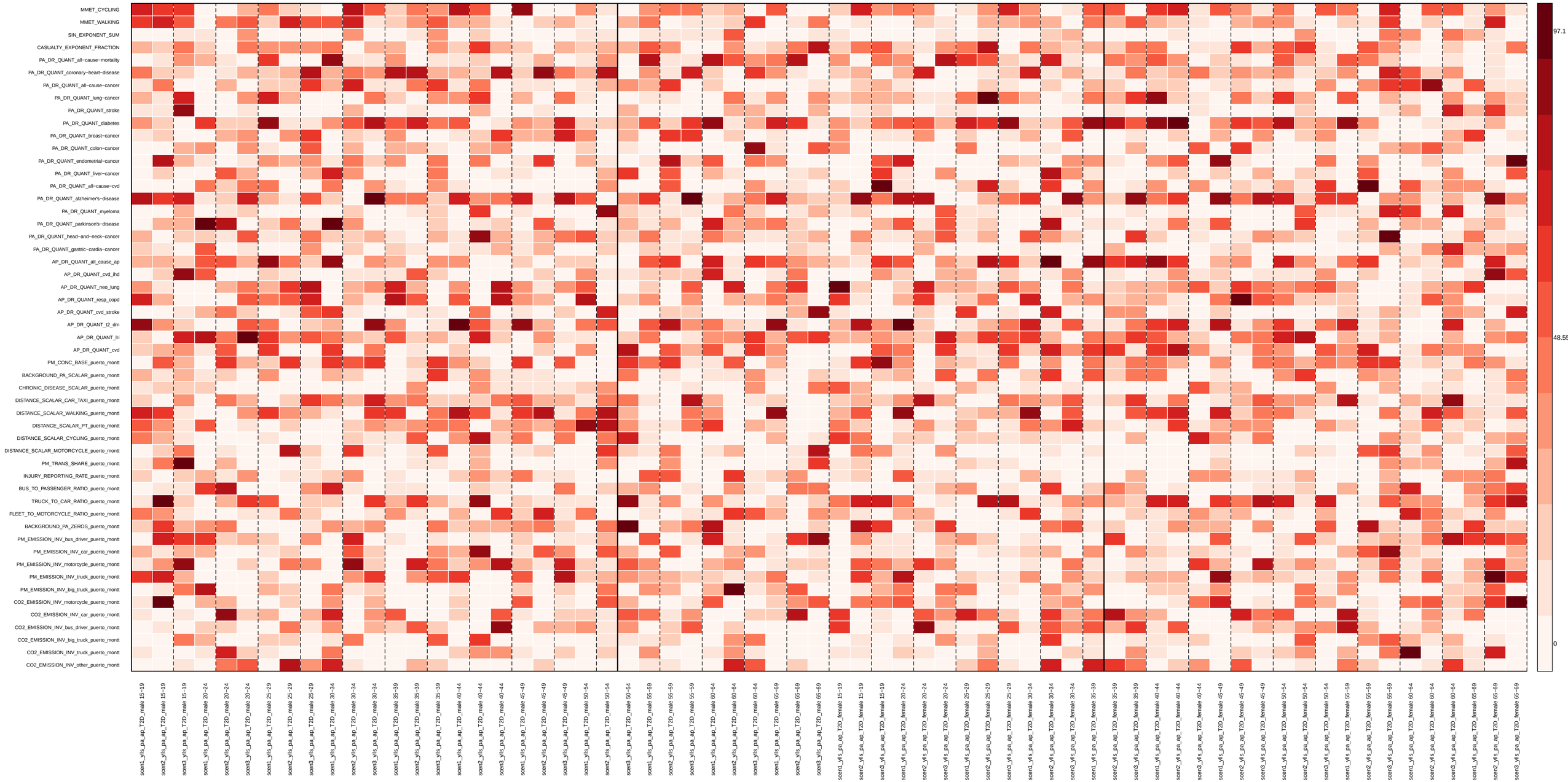
puerto_montt − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



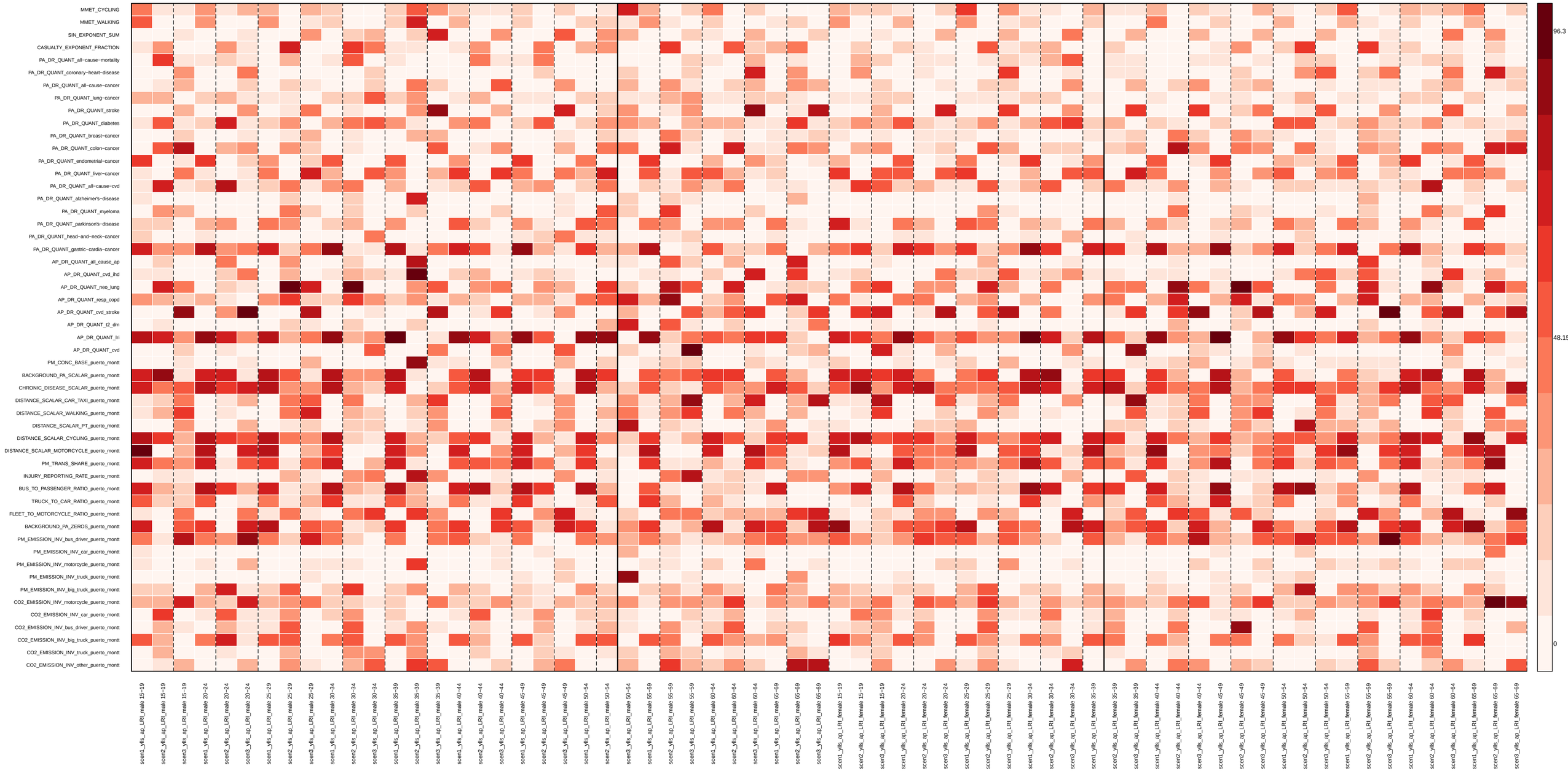
puerto_montt − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



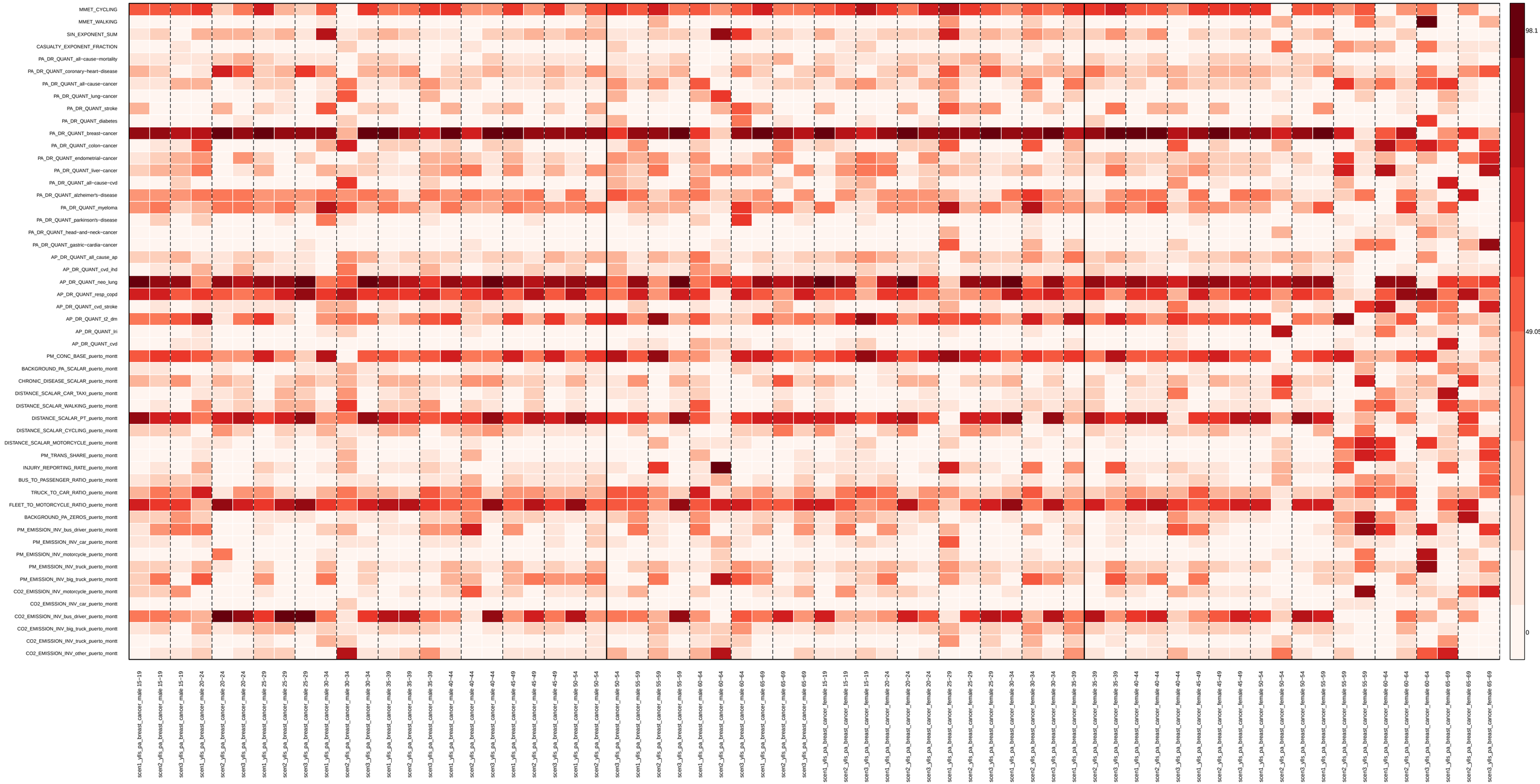
puerto_montt − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



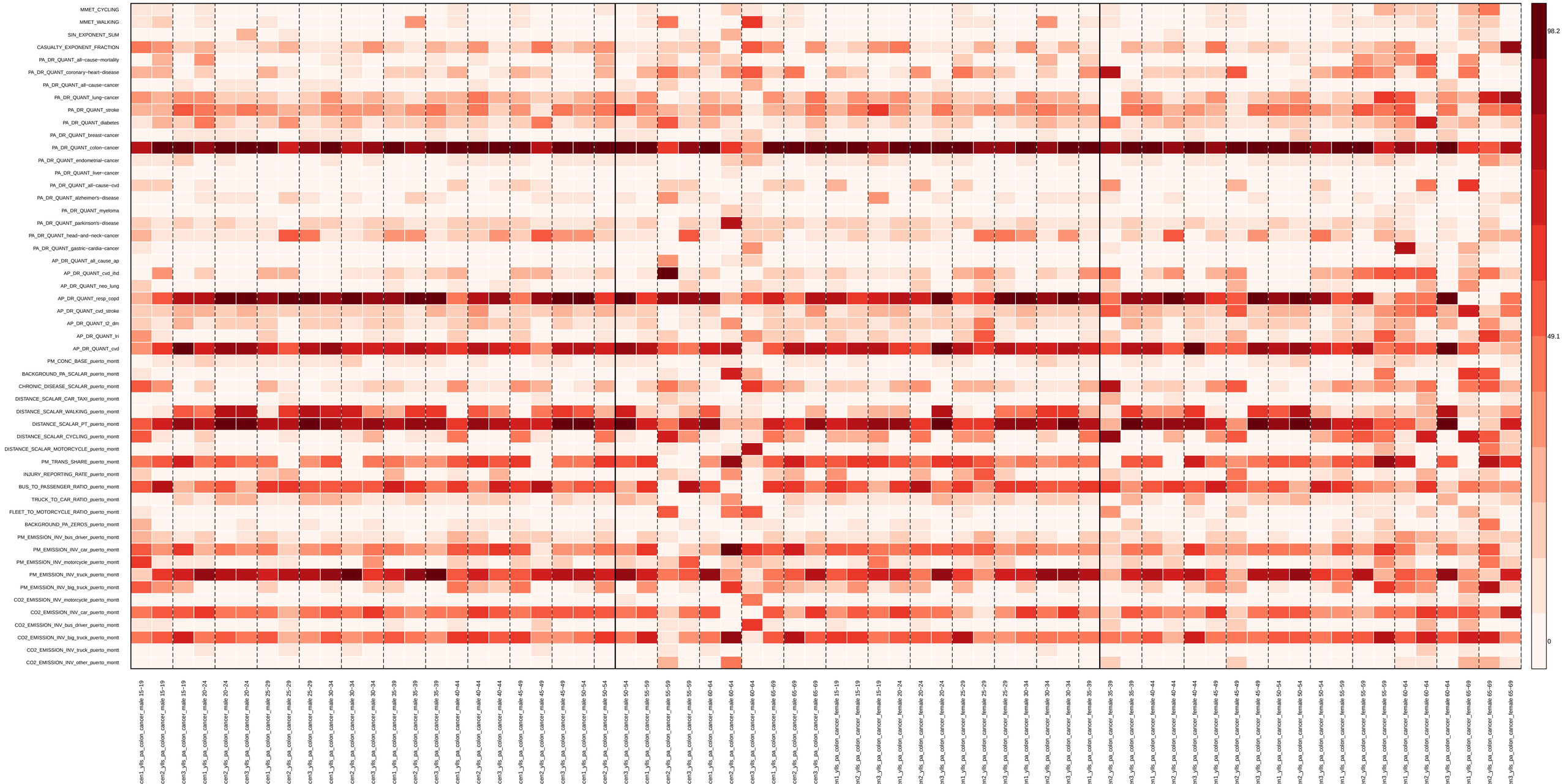
puerto_montt − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



puerto_montt − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

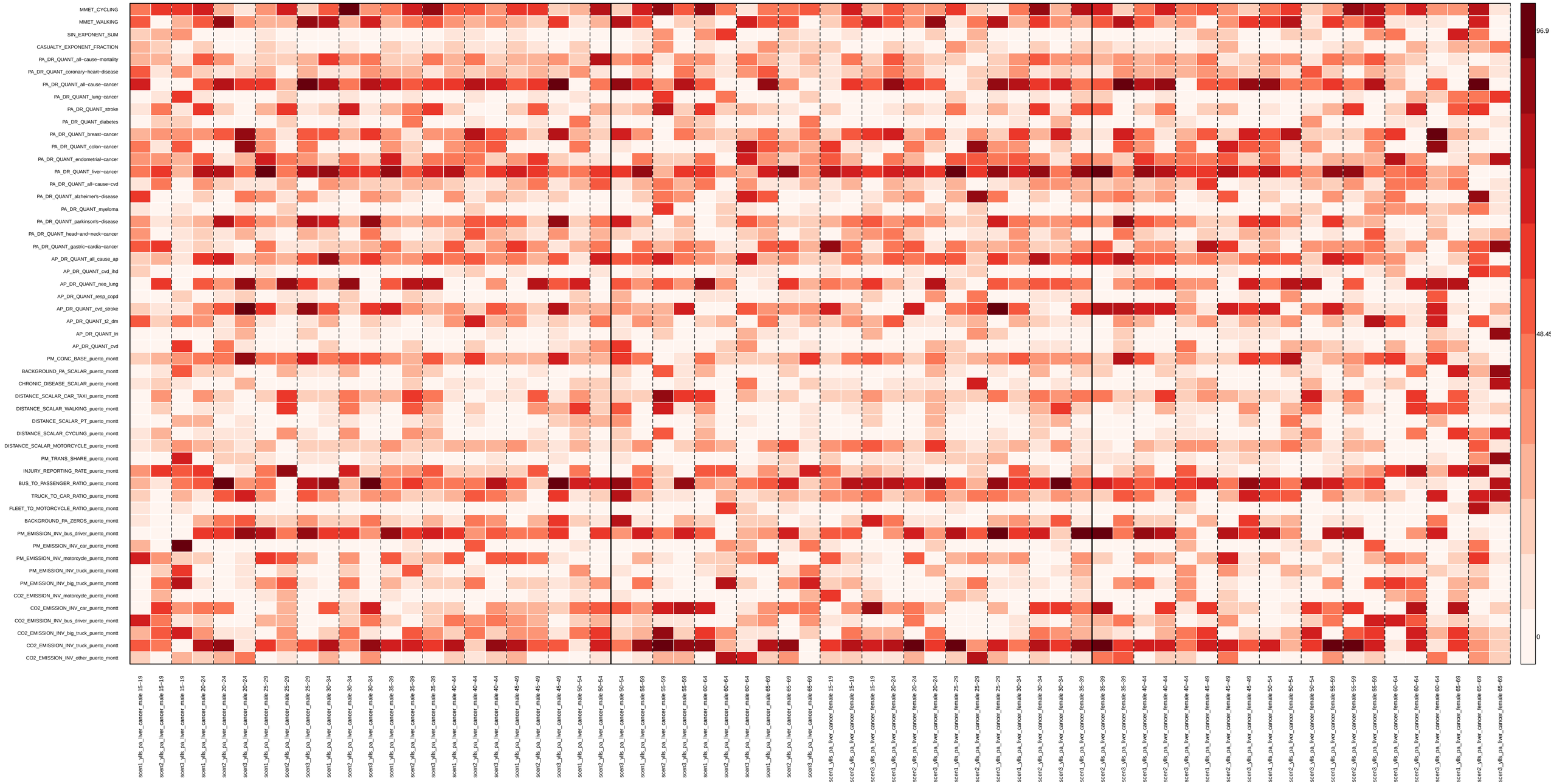


puerto_montt − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

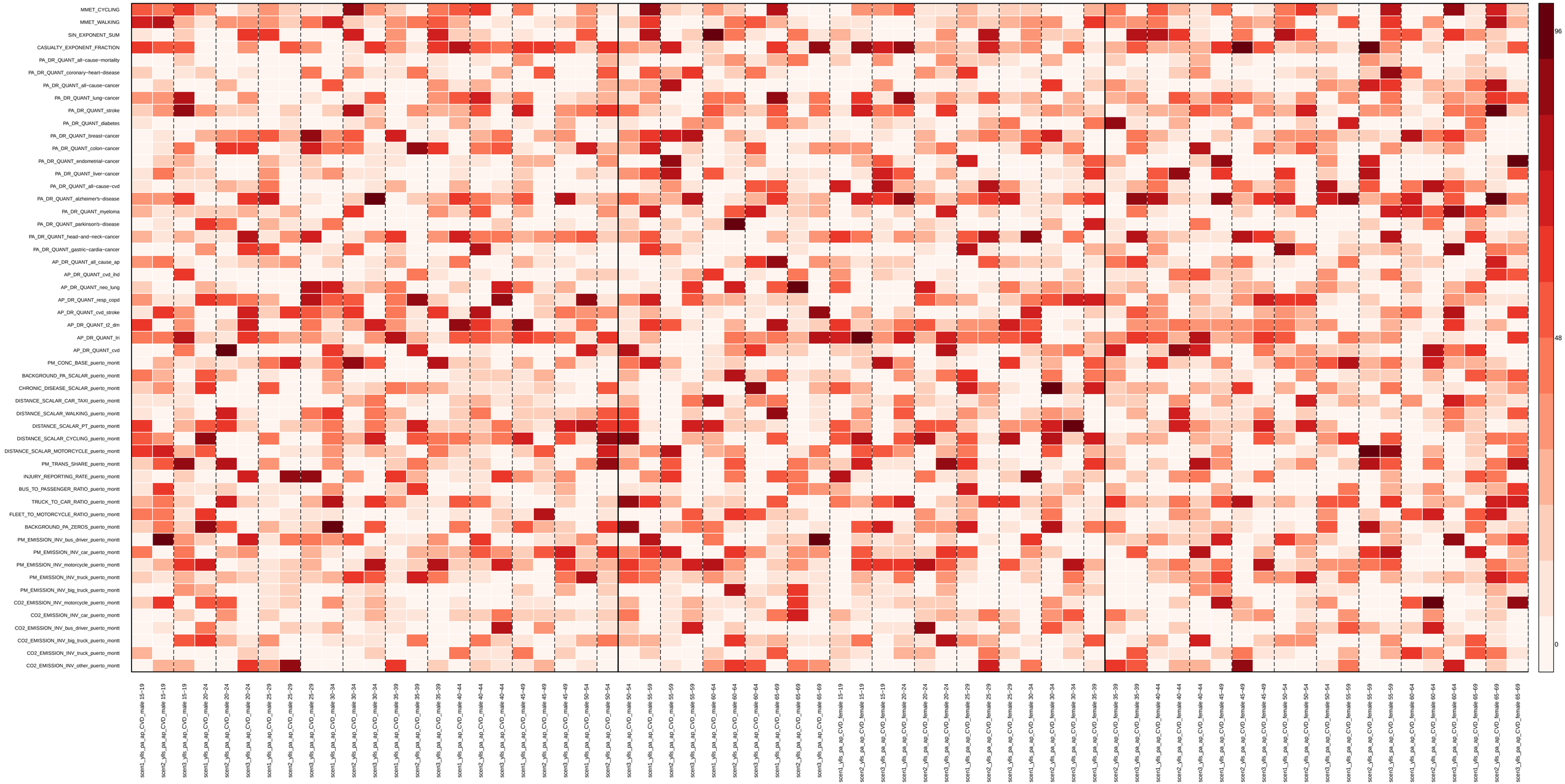


puerto_montt − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

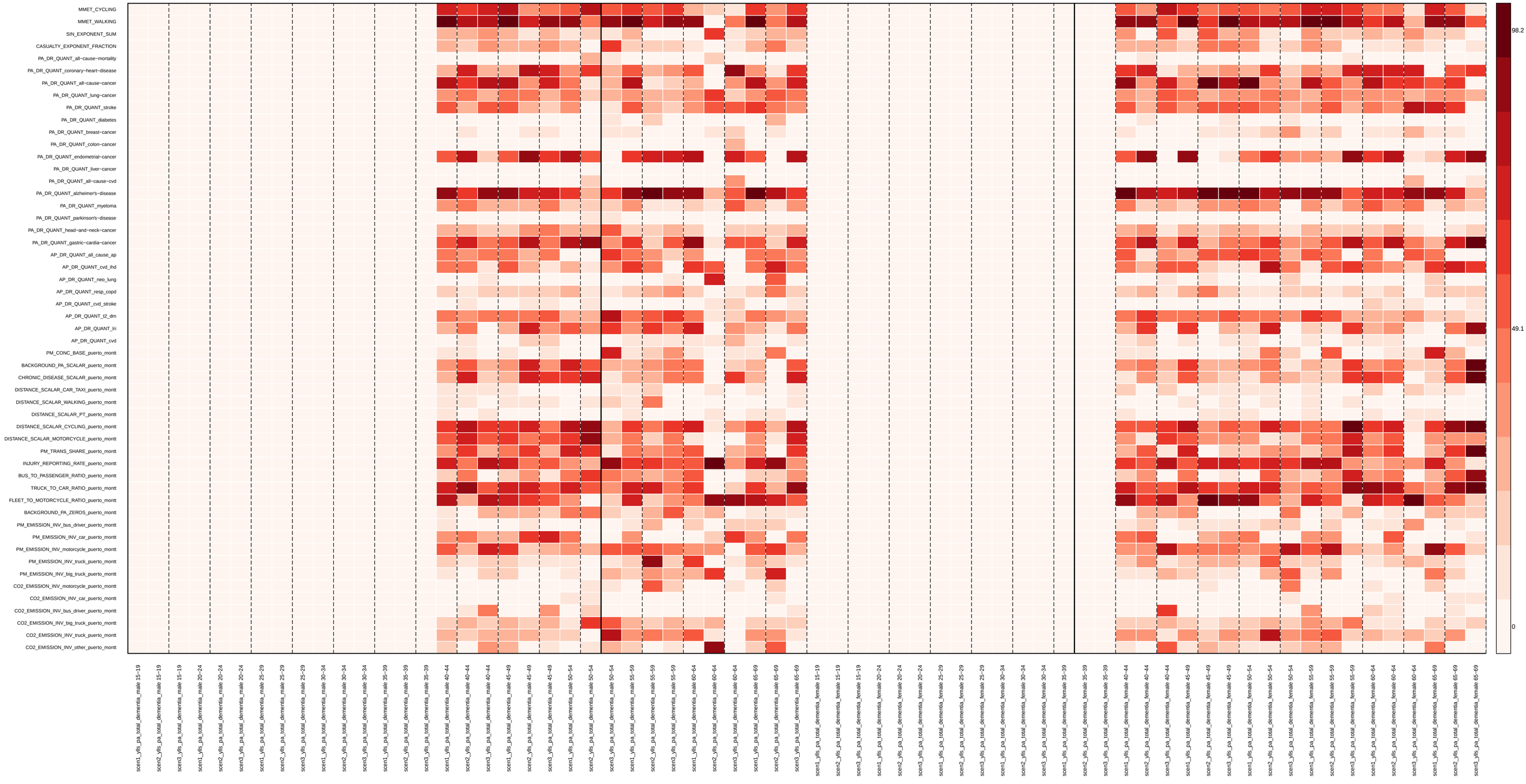
puerto_montt − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



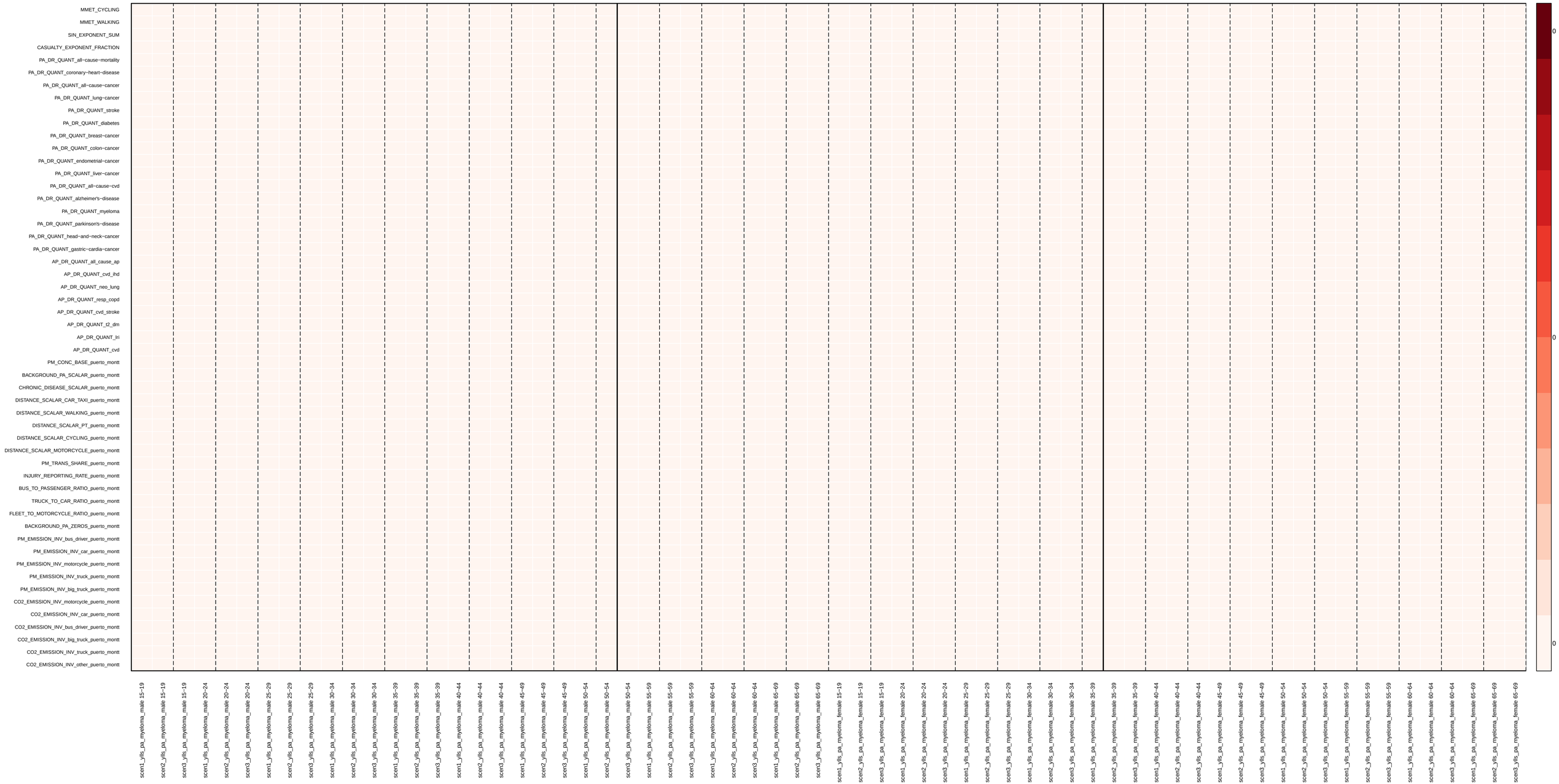
puerto_montt − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



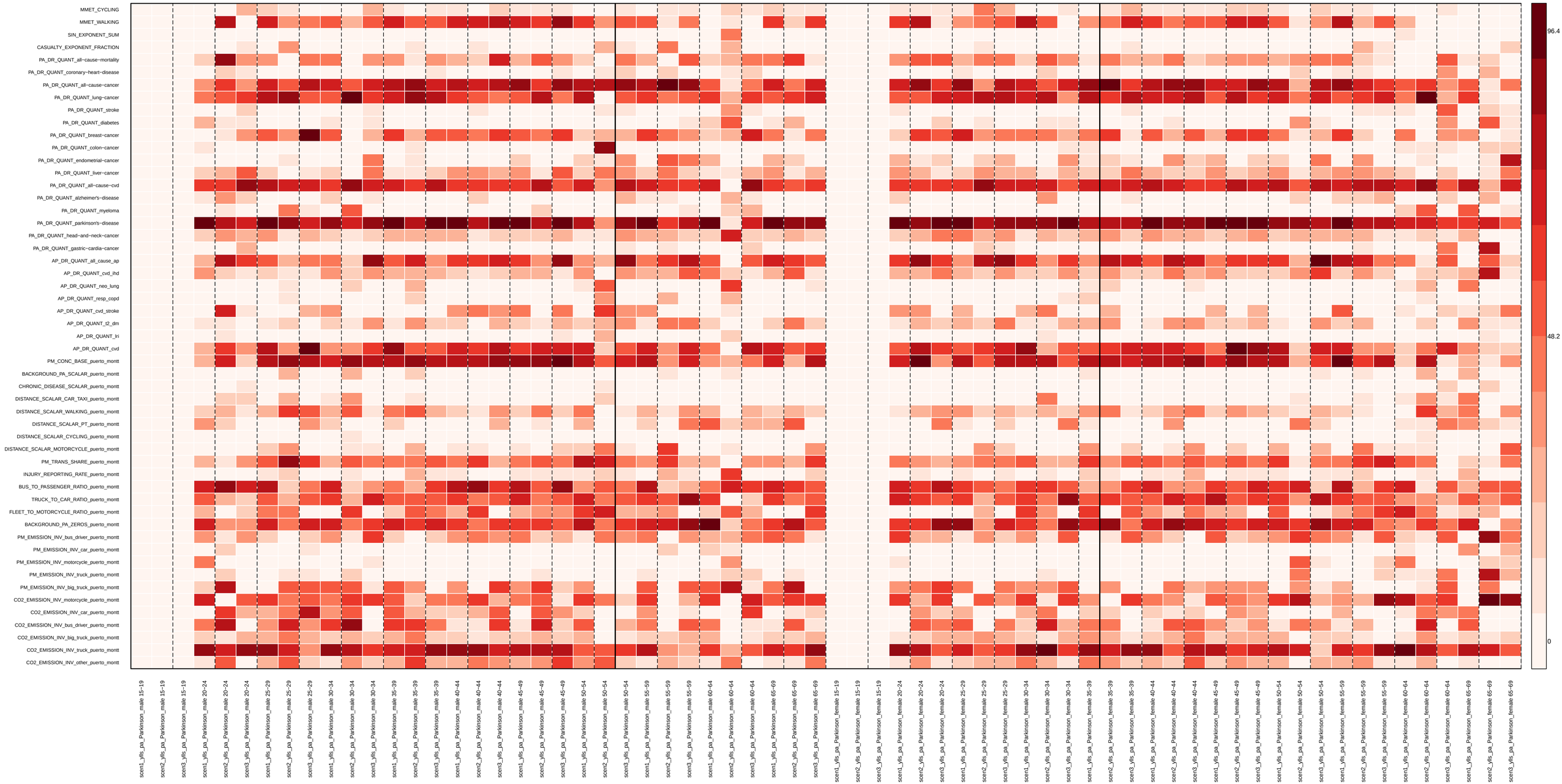
puerto_montt − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



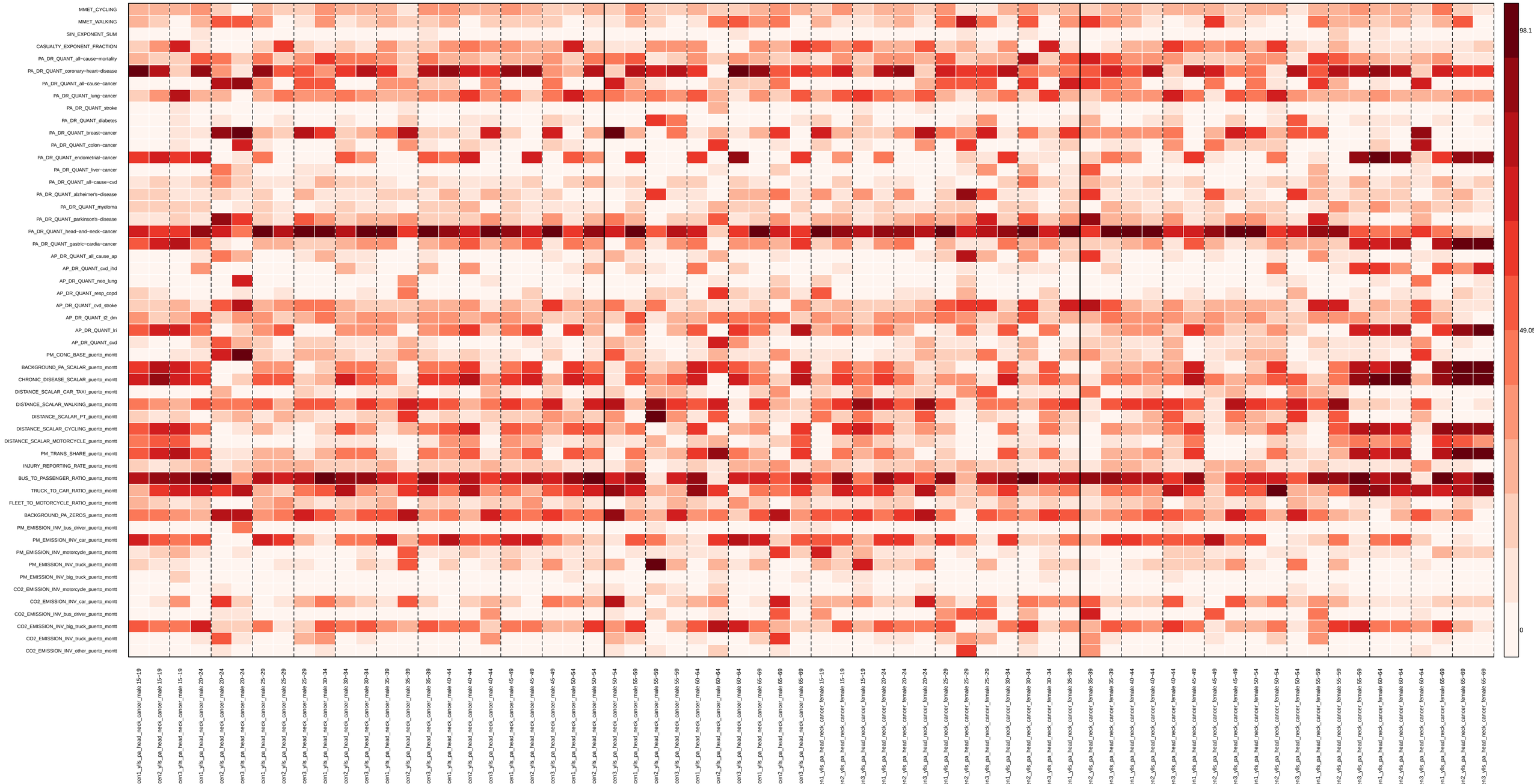
puerto_montt − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



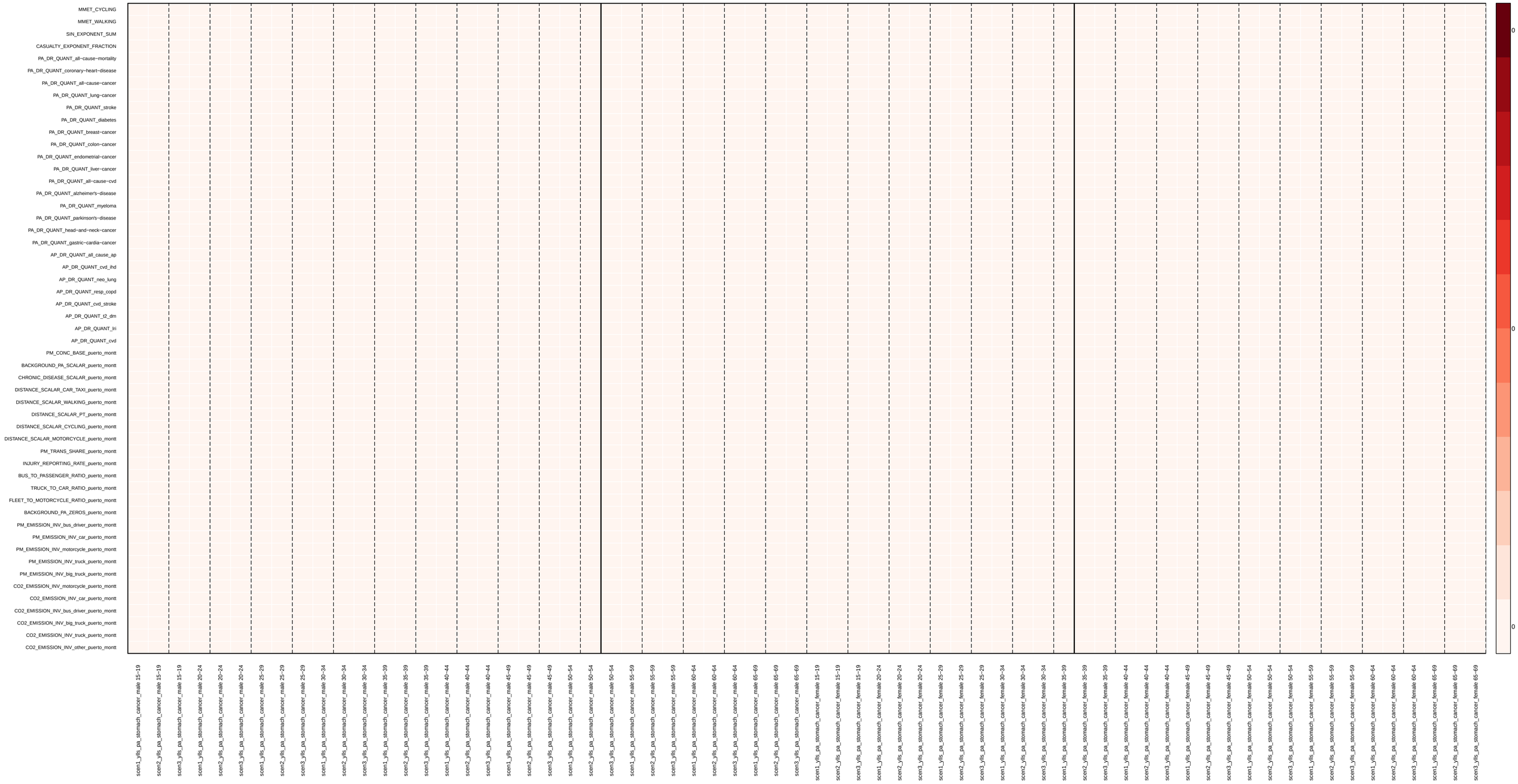
puerto_montt − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



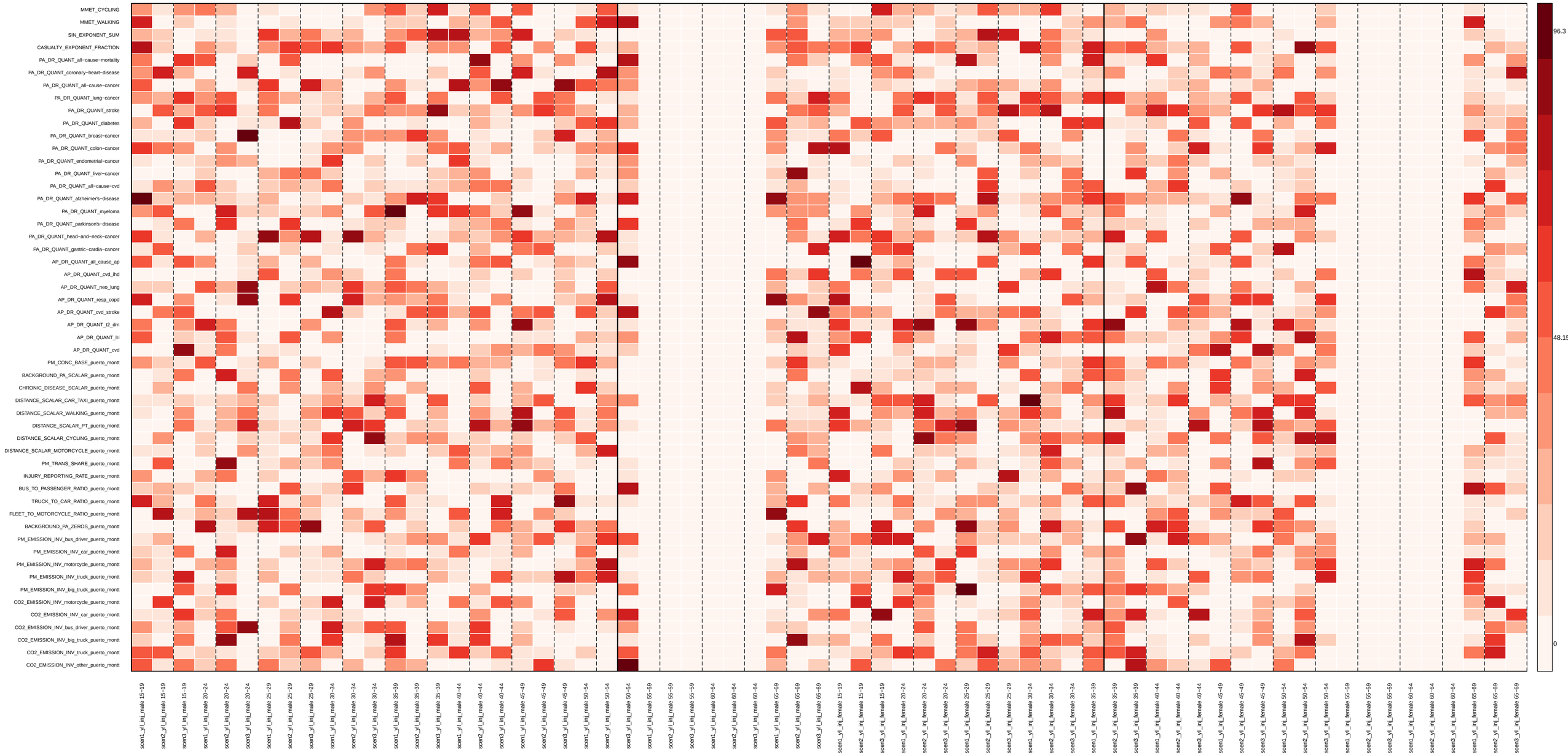
puerto_montt − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



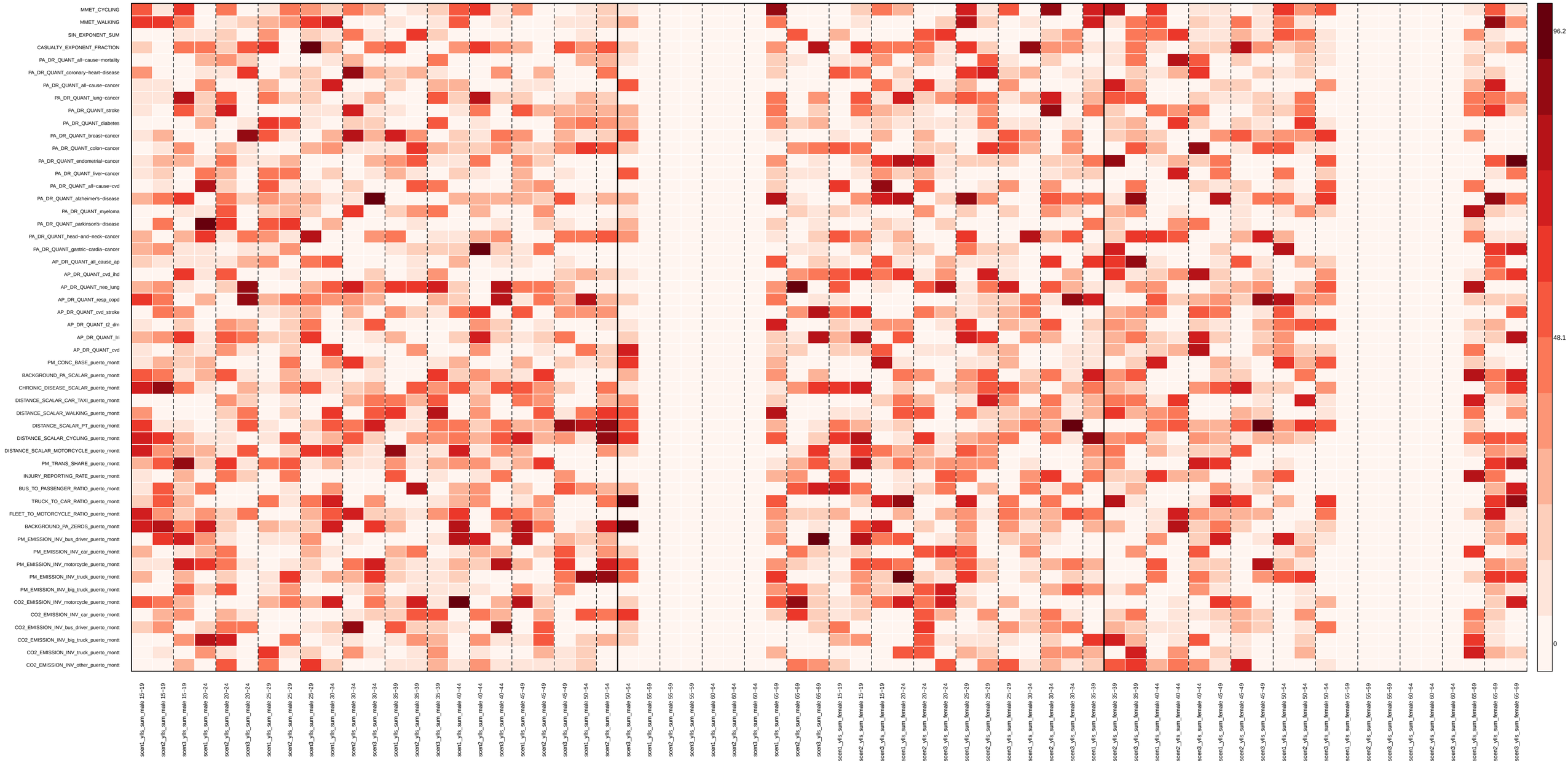
puerto_montt − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



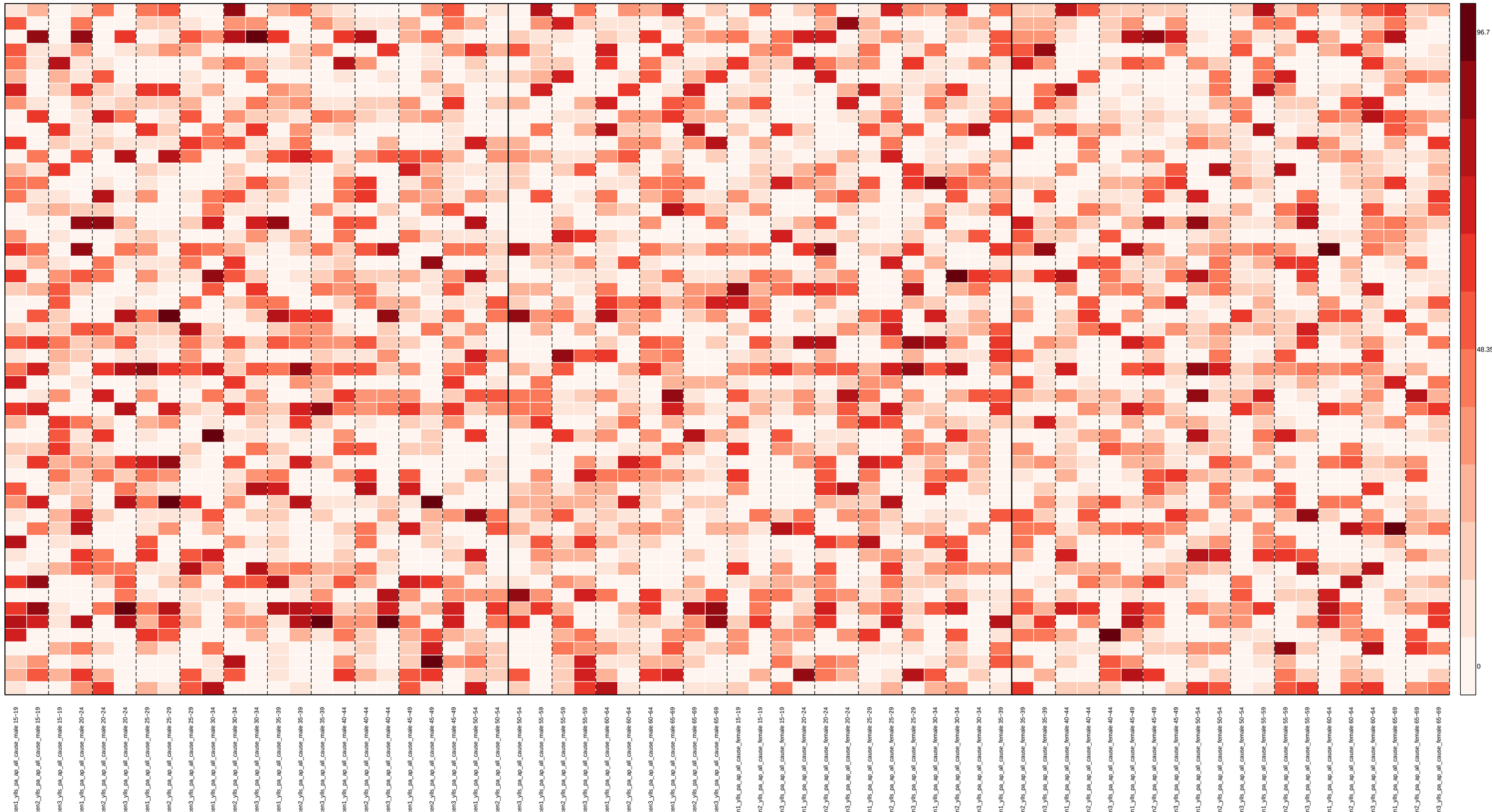
puerto_montt − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



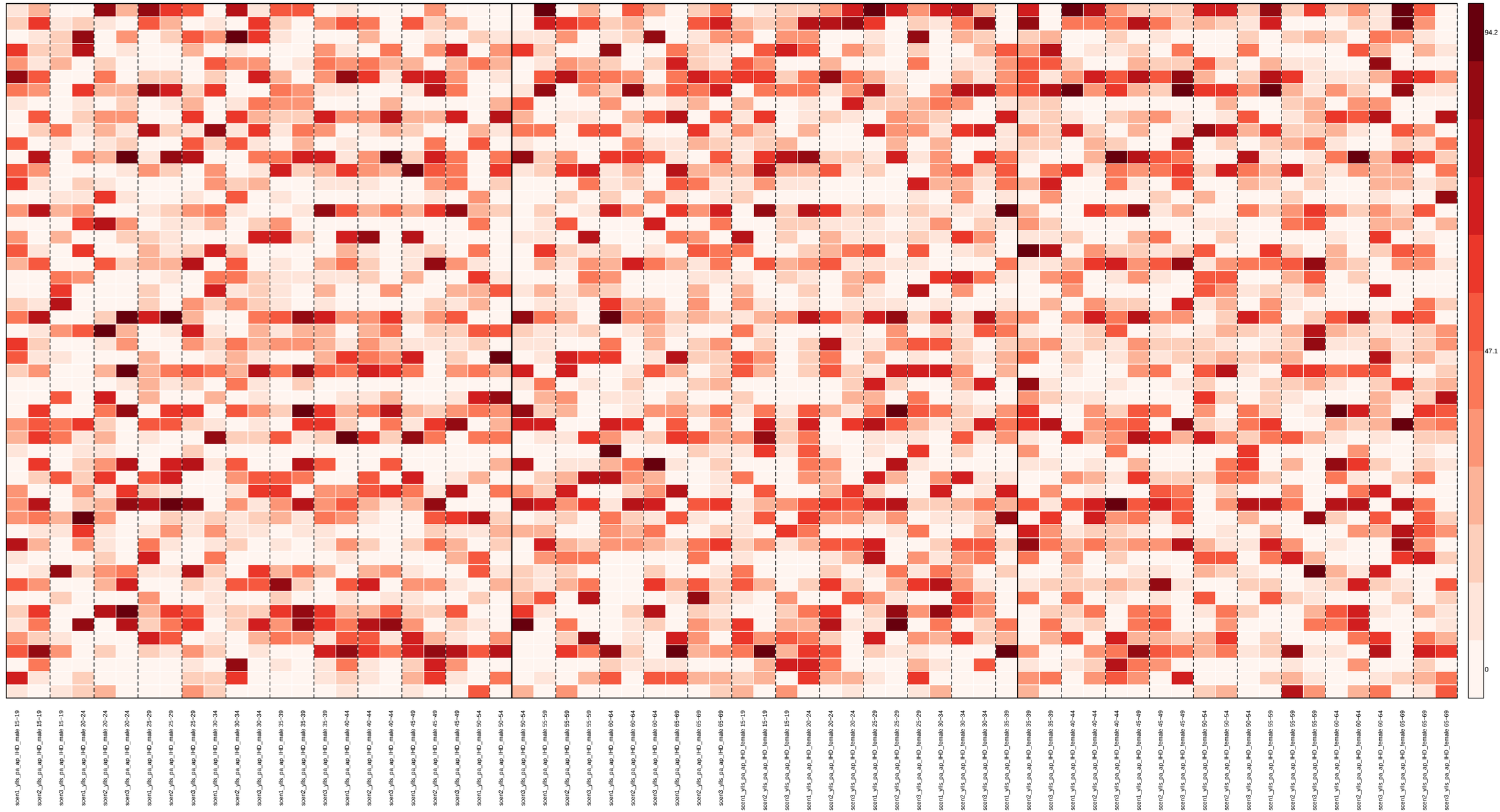
puerto_montt − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



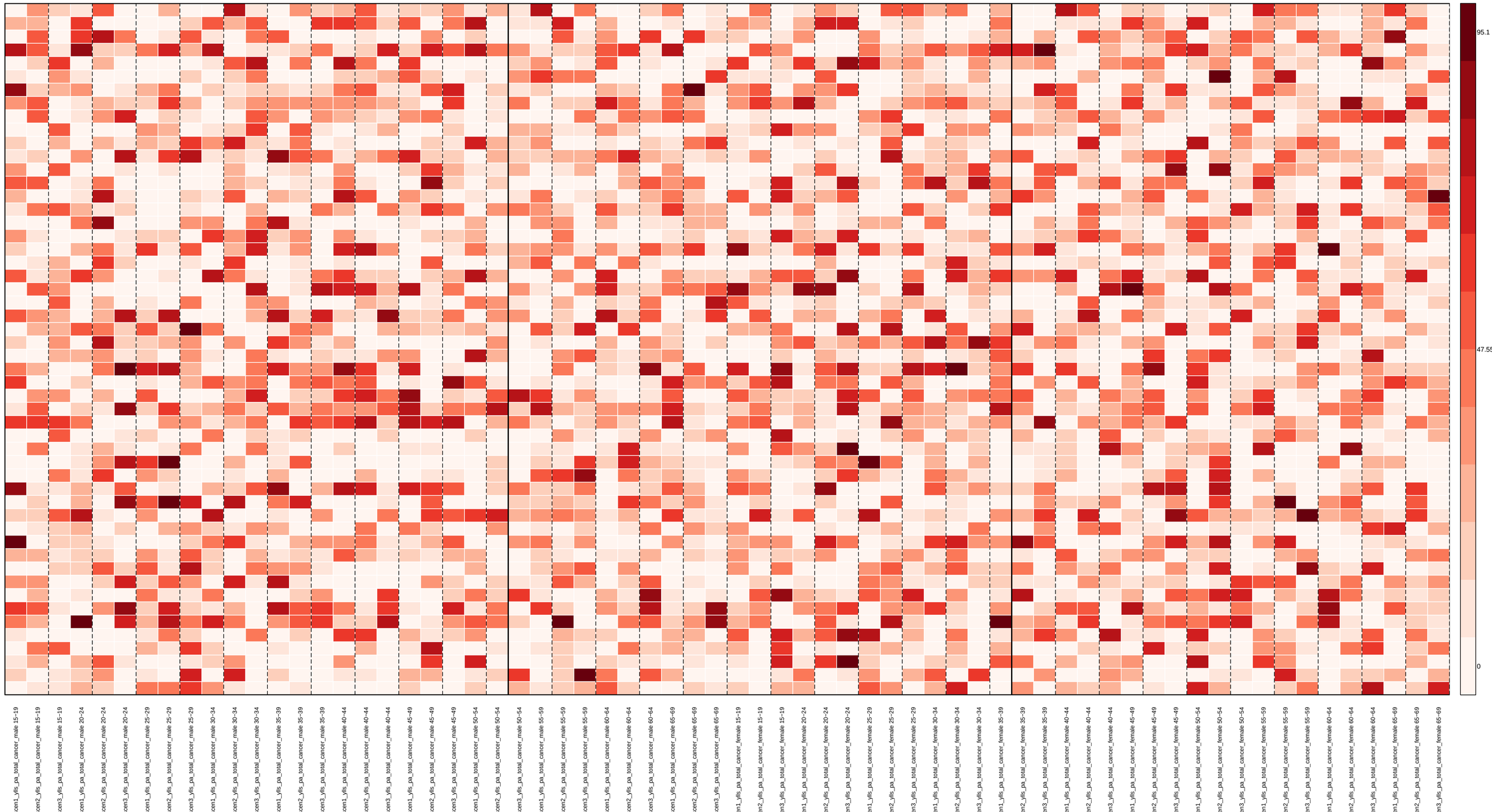
san_antonio − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



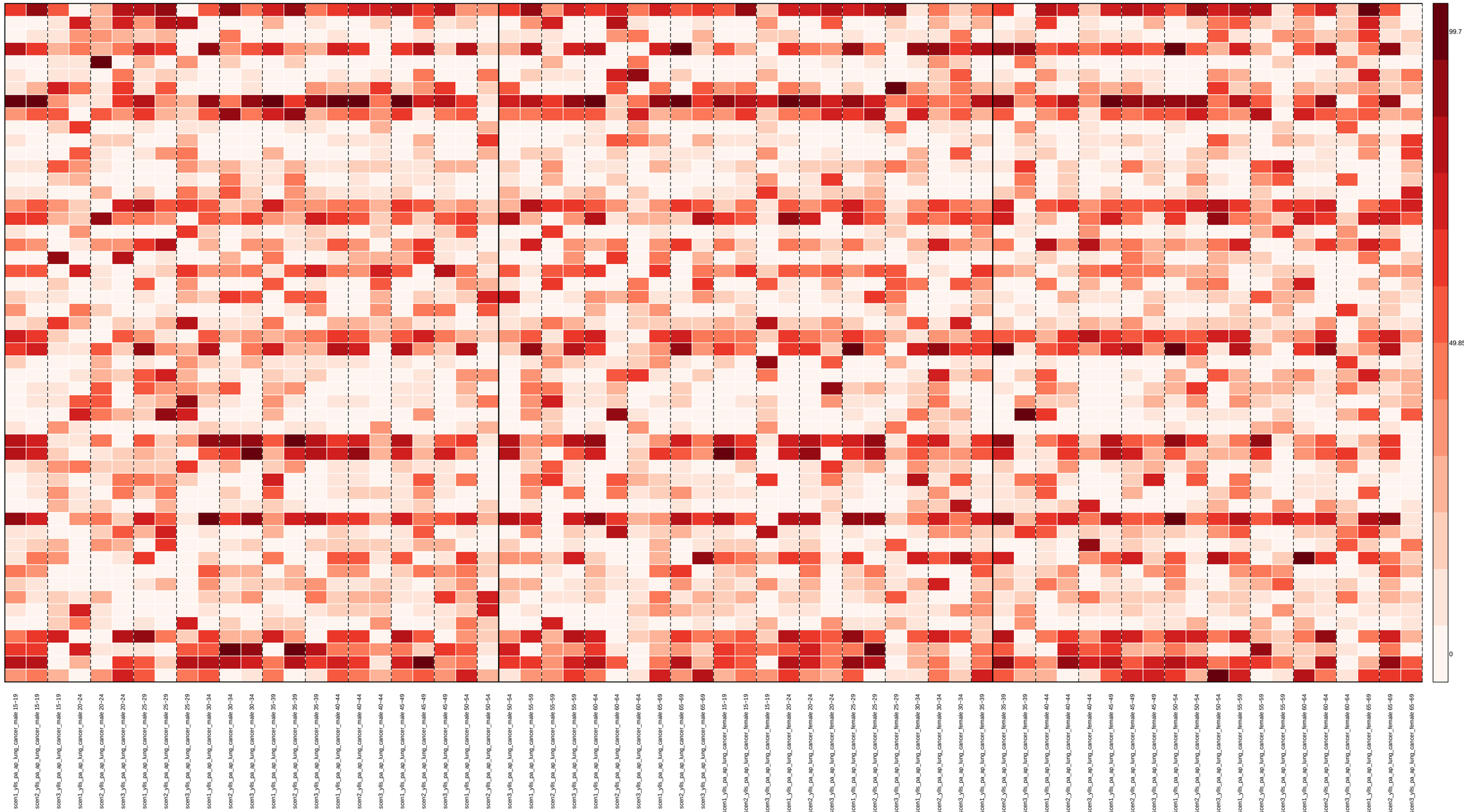
san_antonio − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



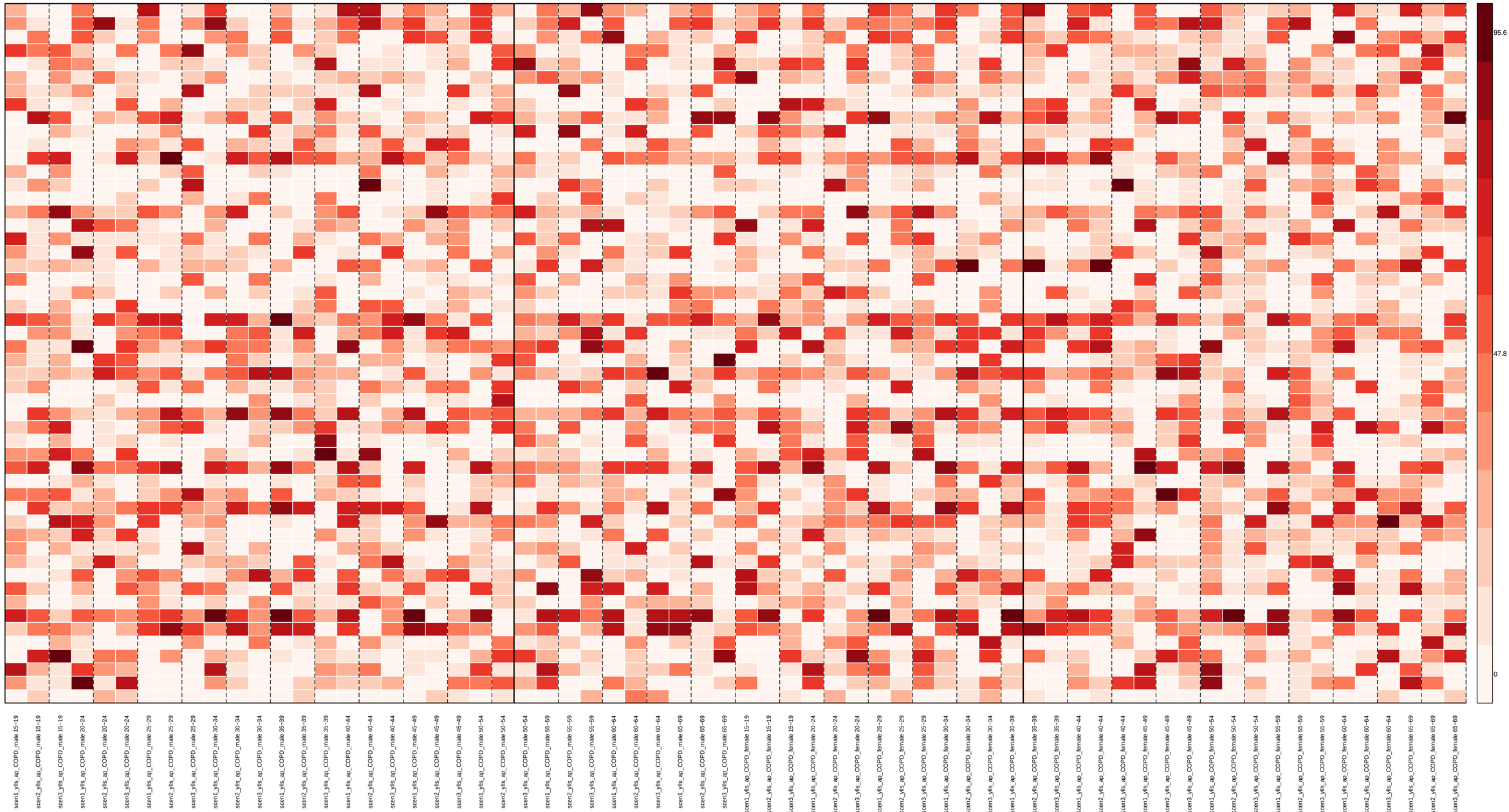
san_antonio − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



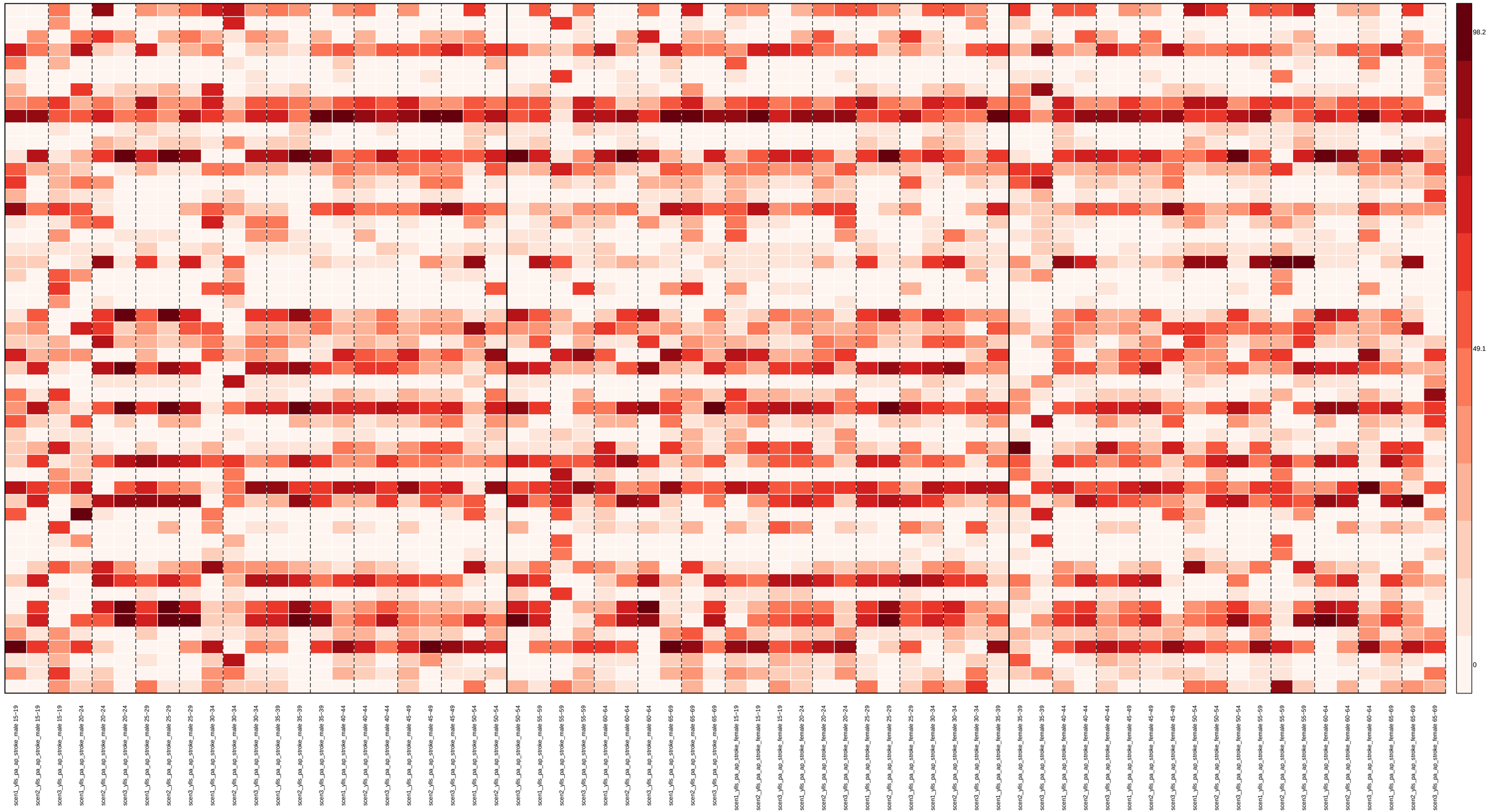
san_antonio − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



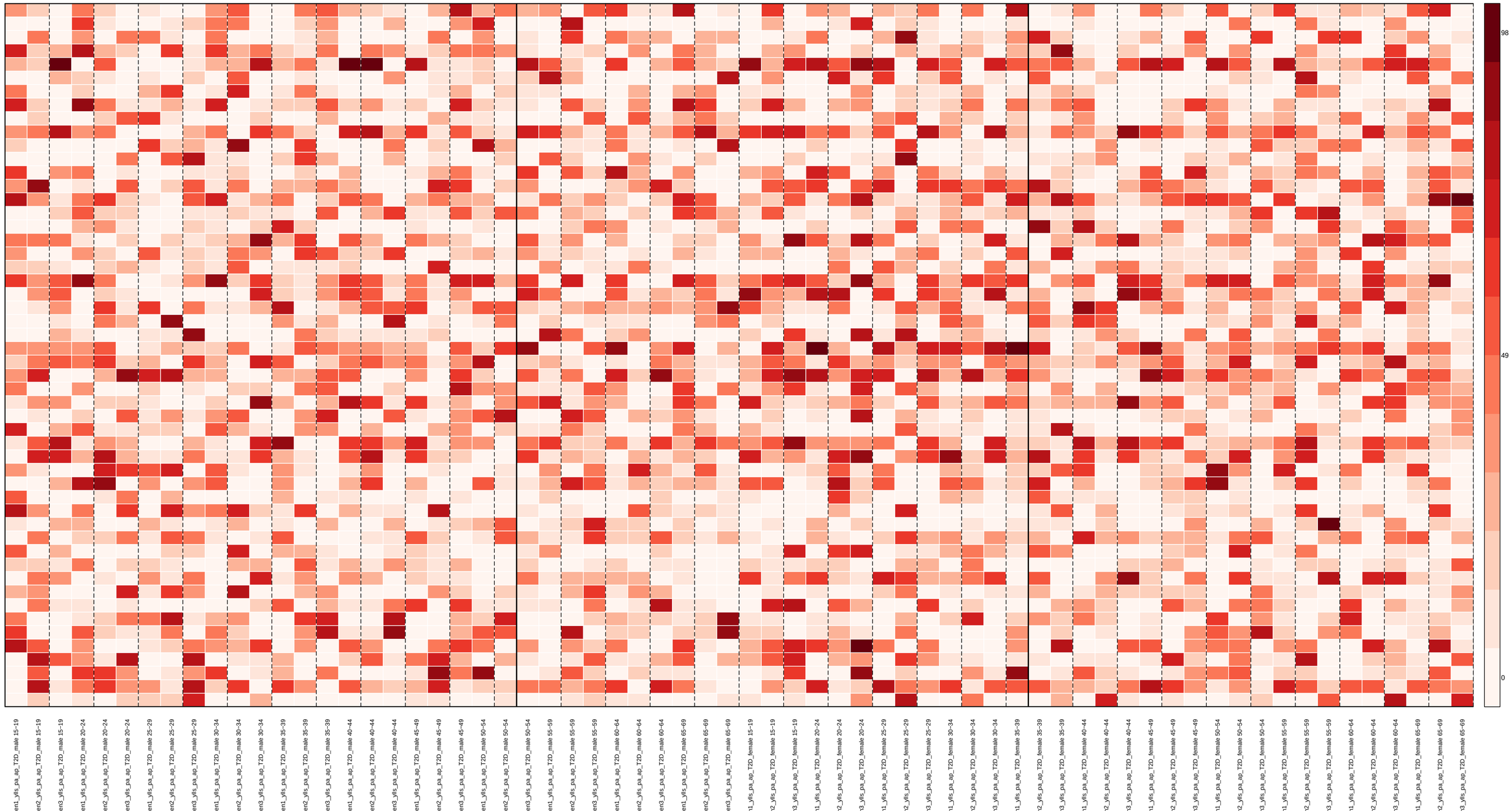
san_antonio − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



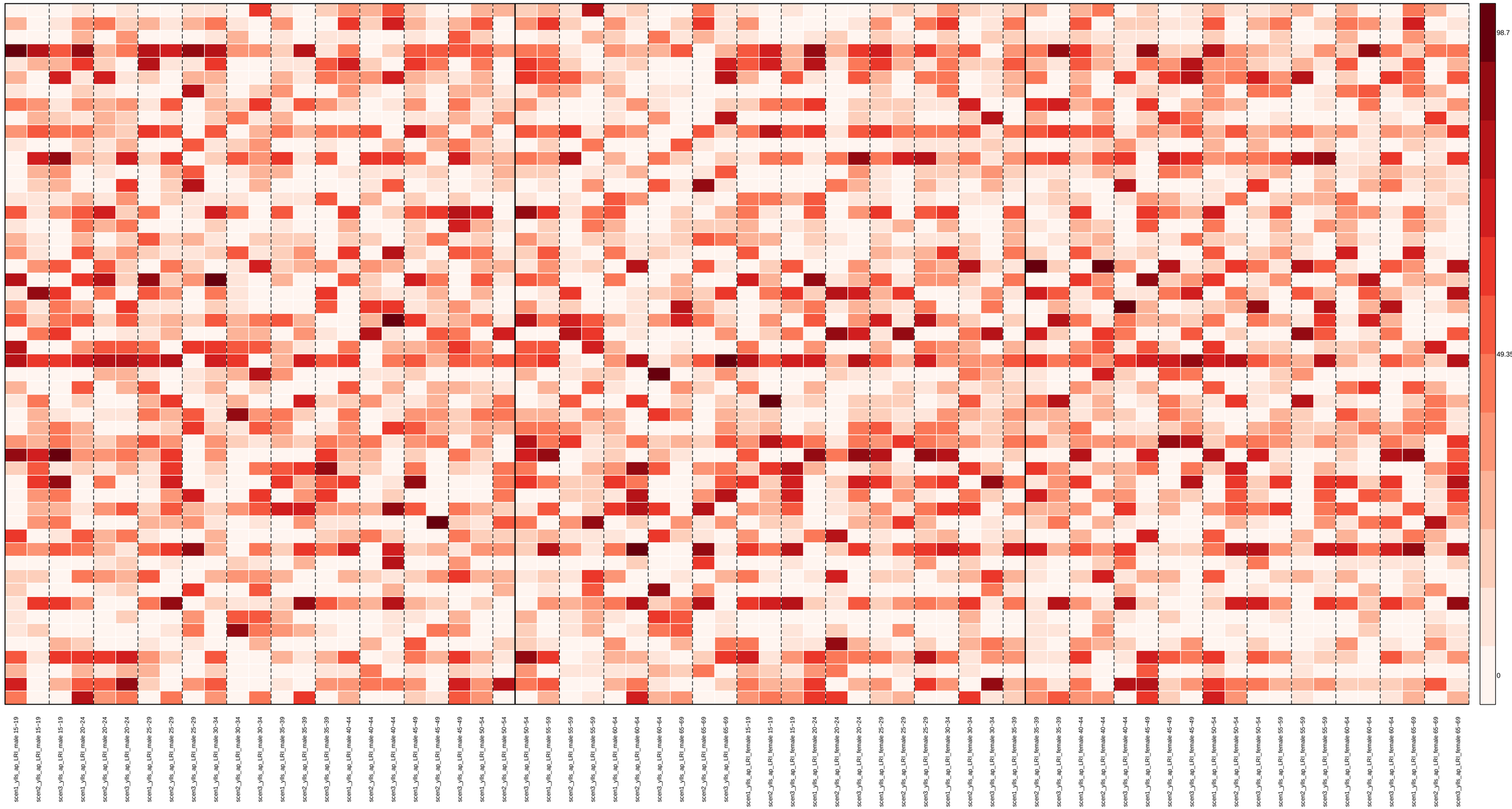
san_antonio − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



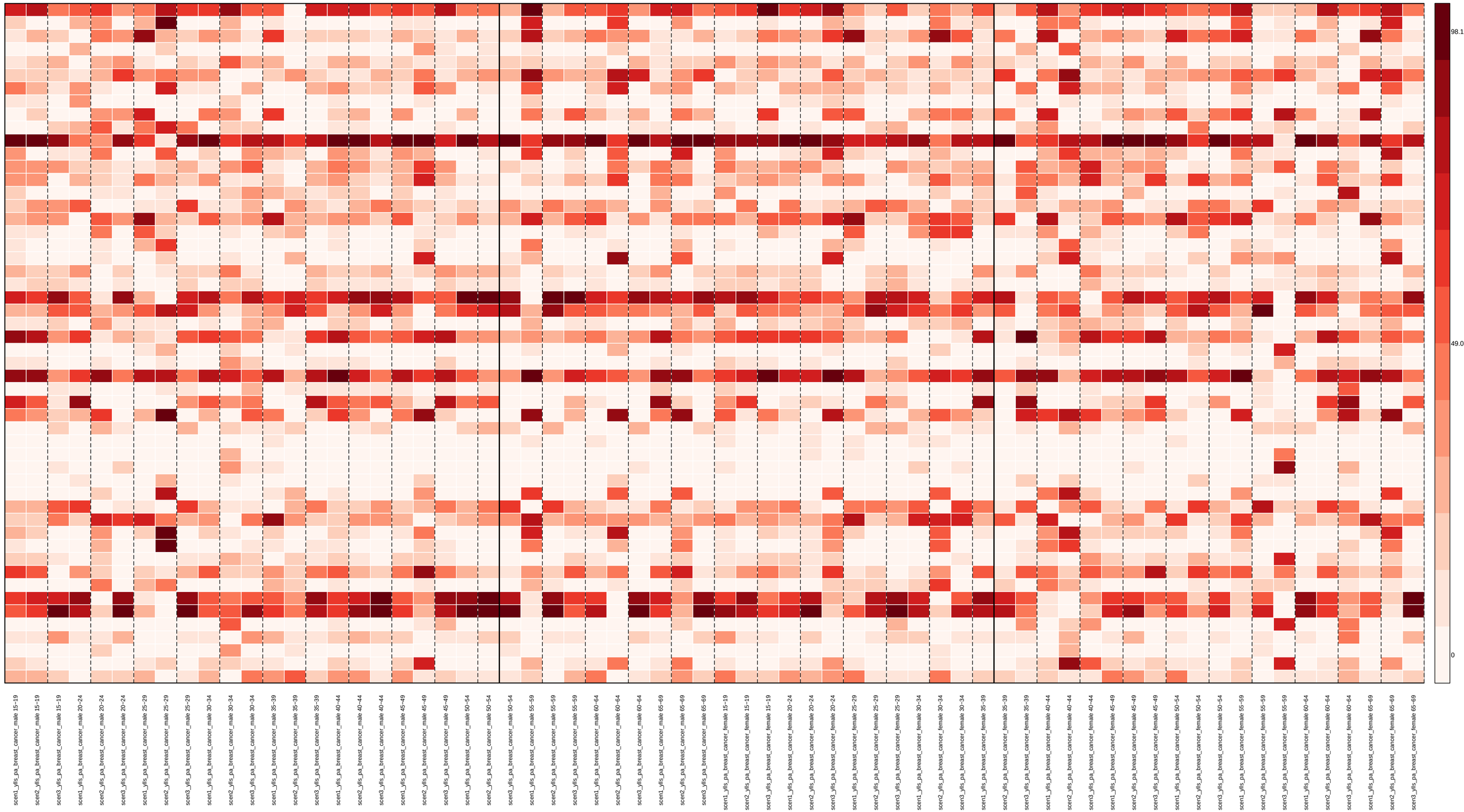
san_antonio − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



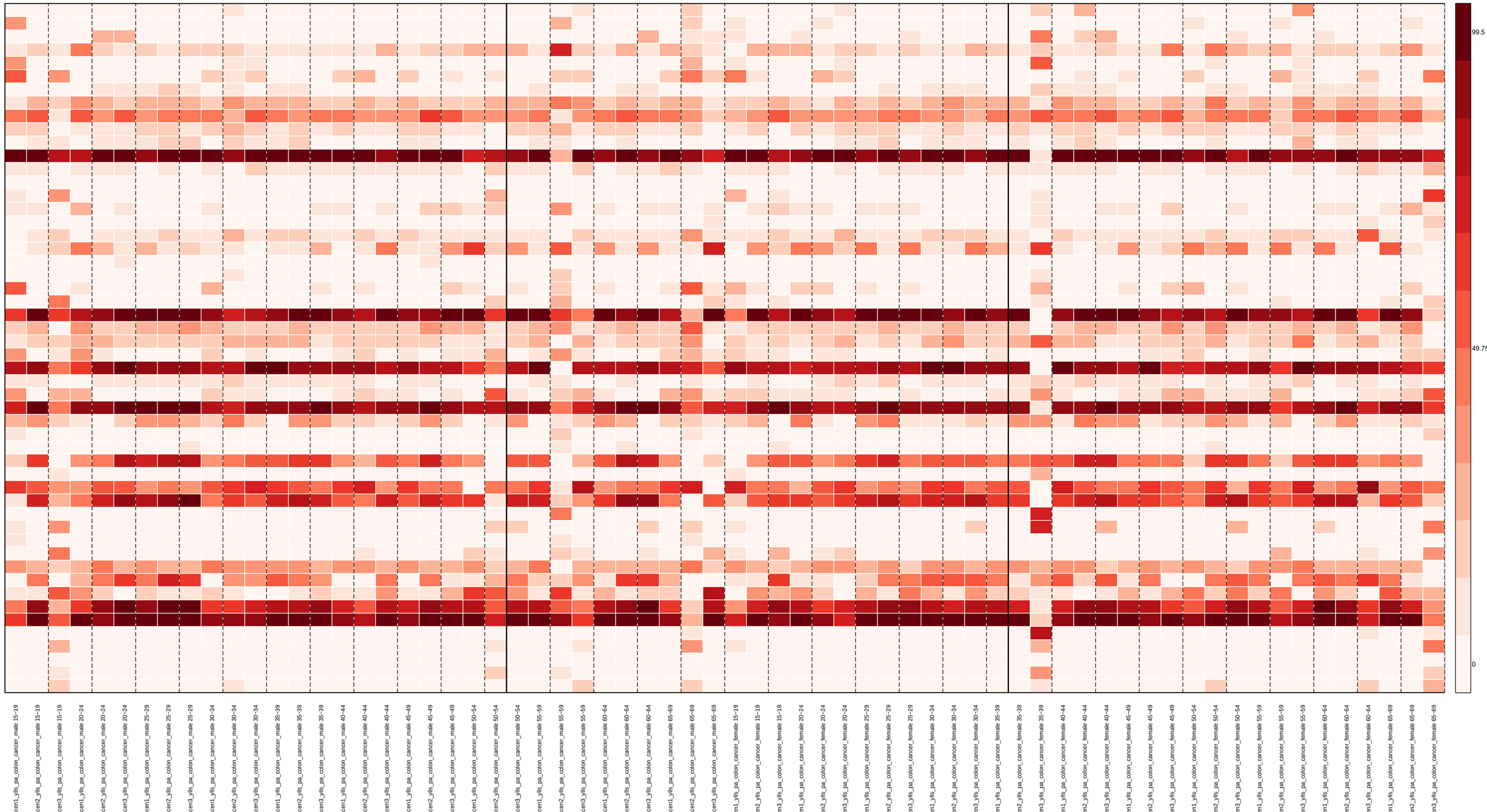
san_antonio − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



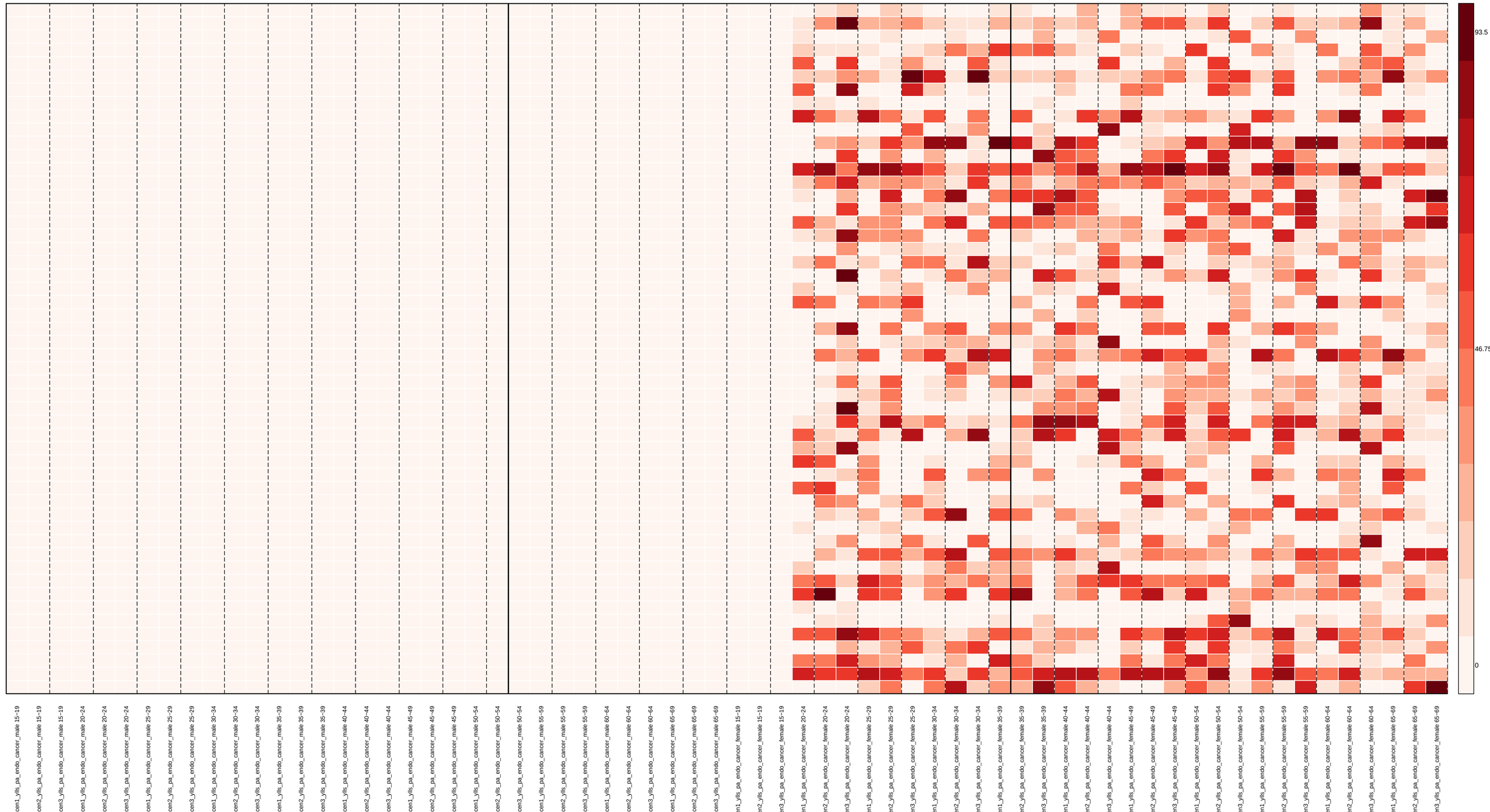
san_antonio − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



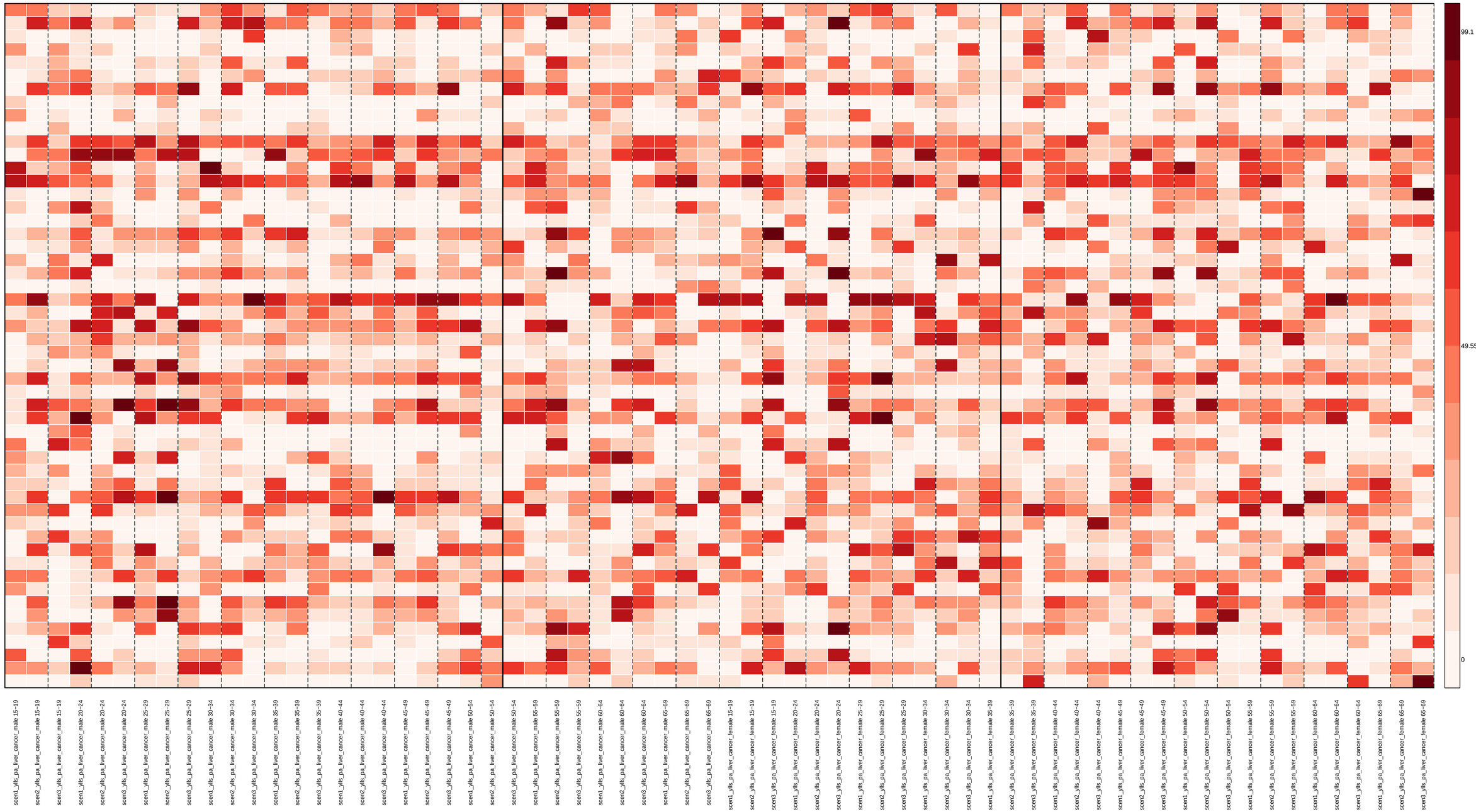
san_antonio − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



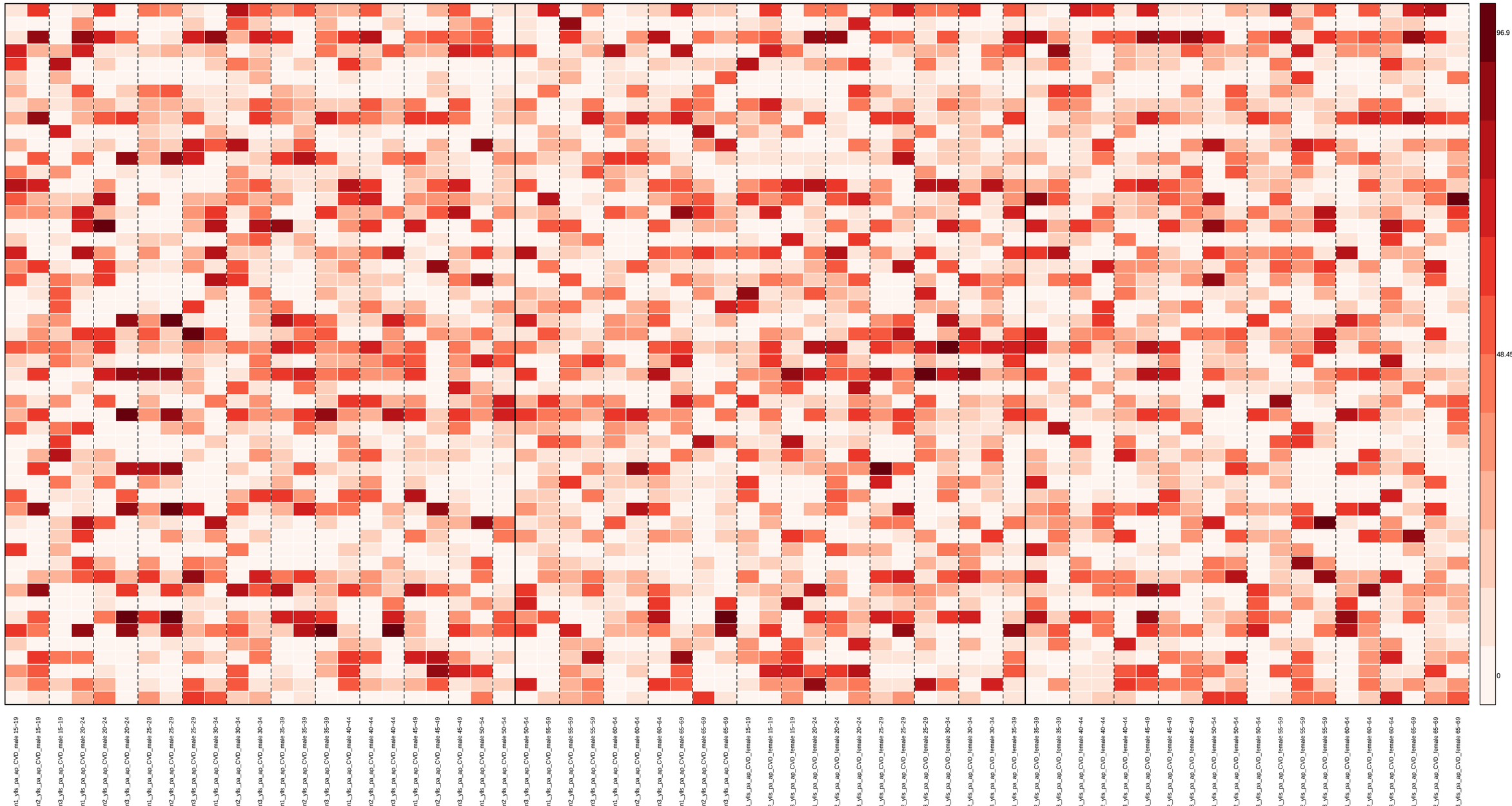
san_antonio − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



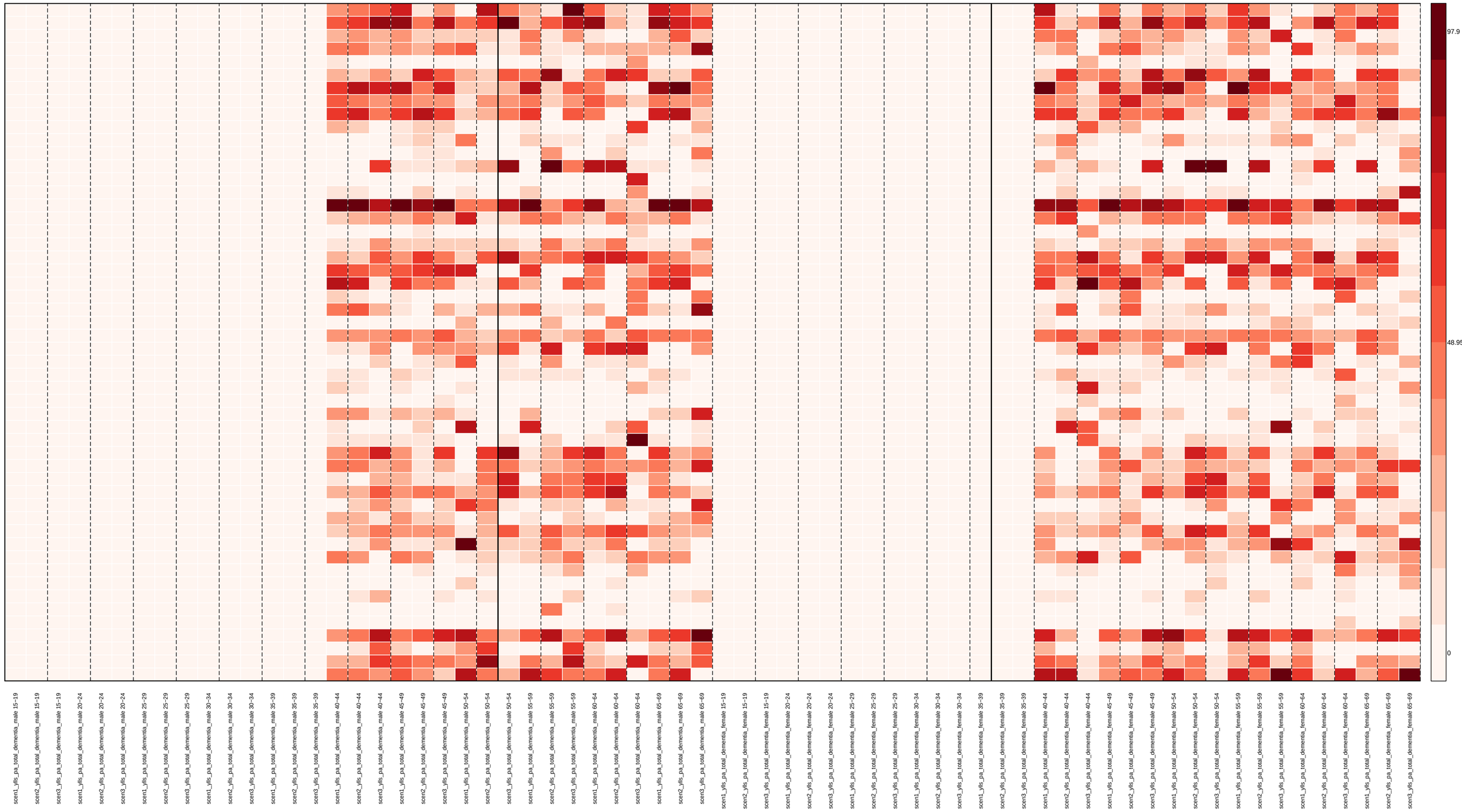
san_antonio − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



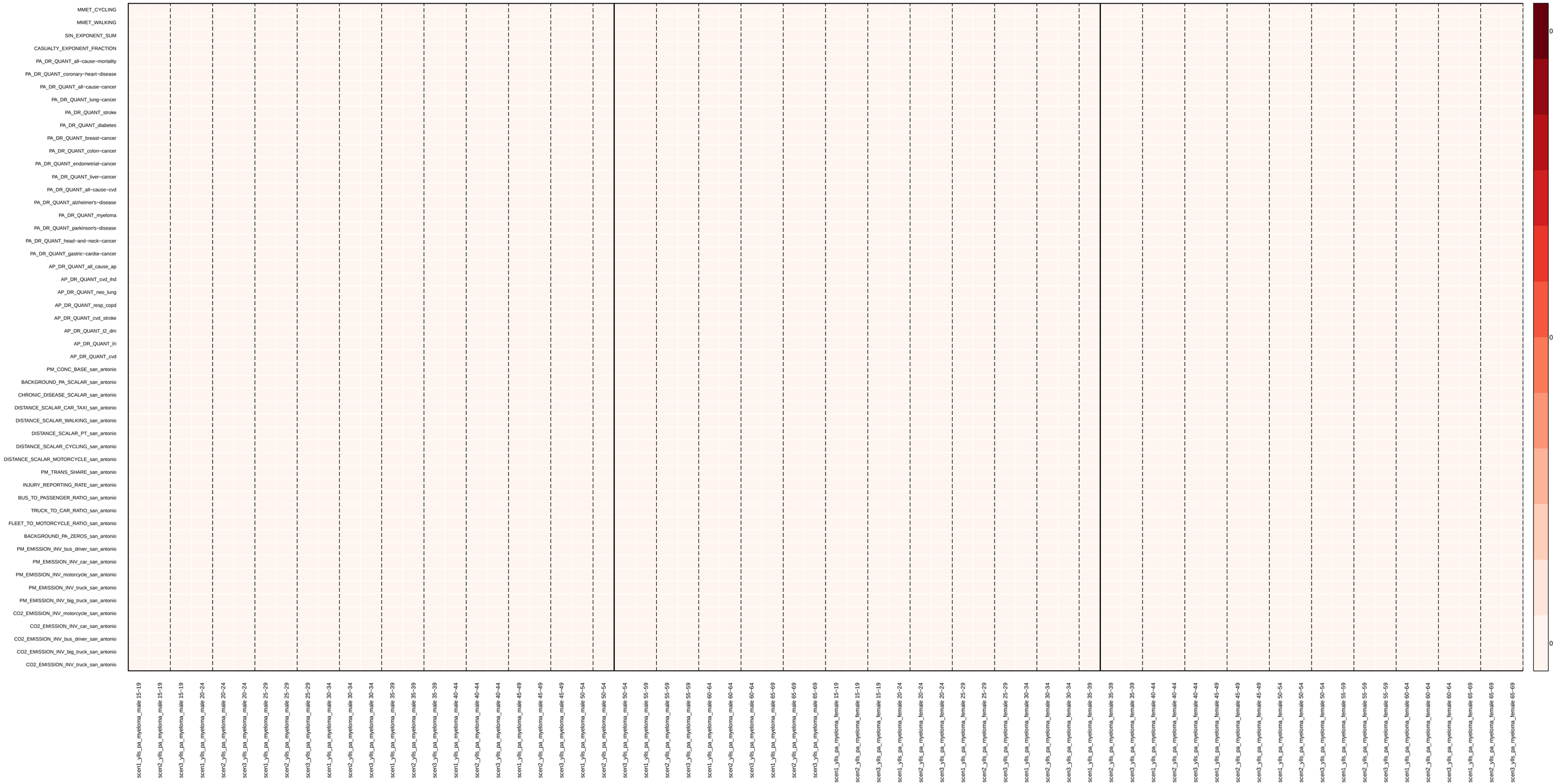
san_antonio − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



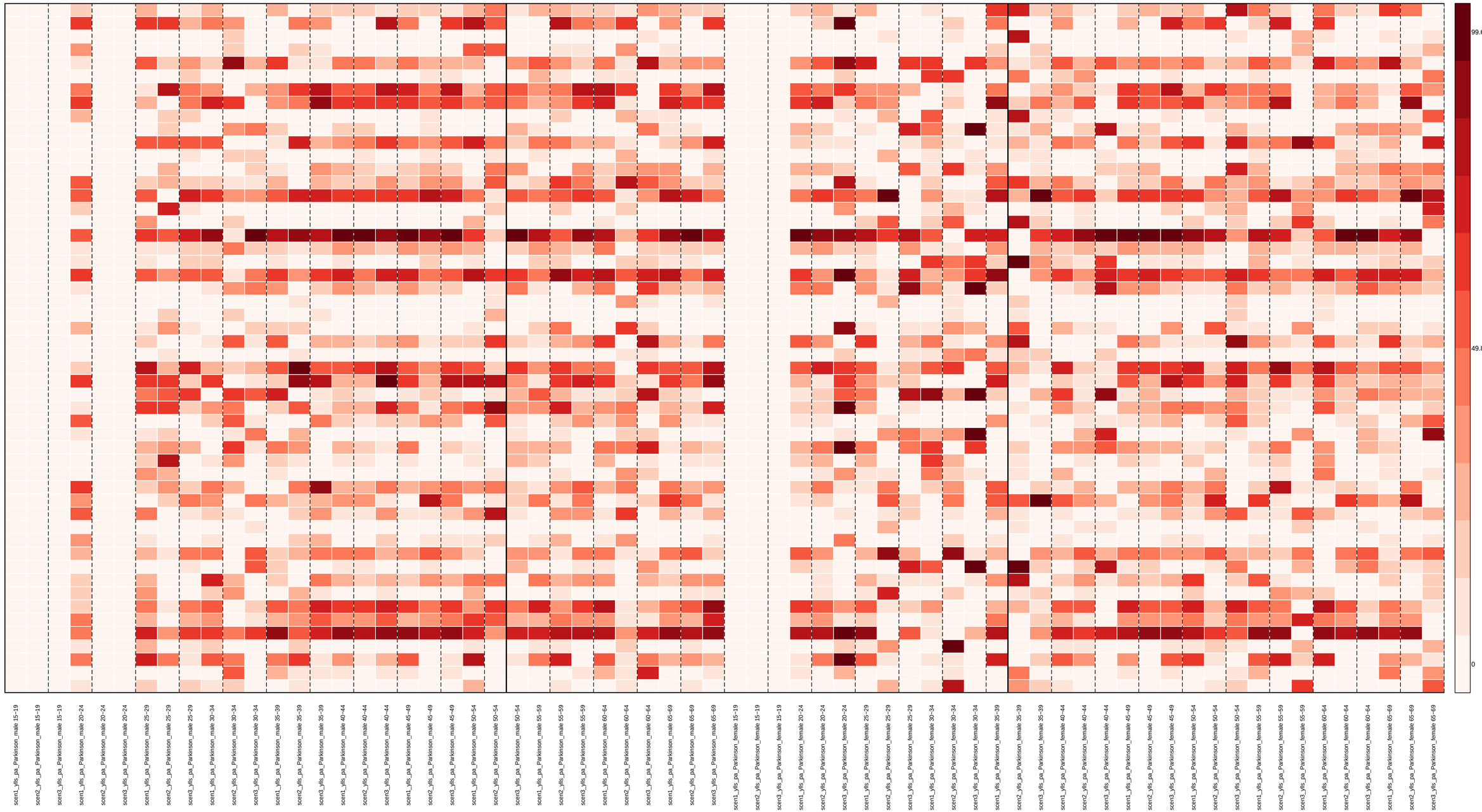
san_antonio − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



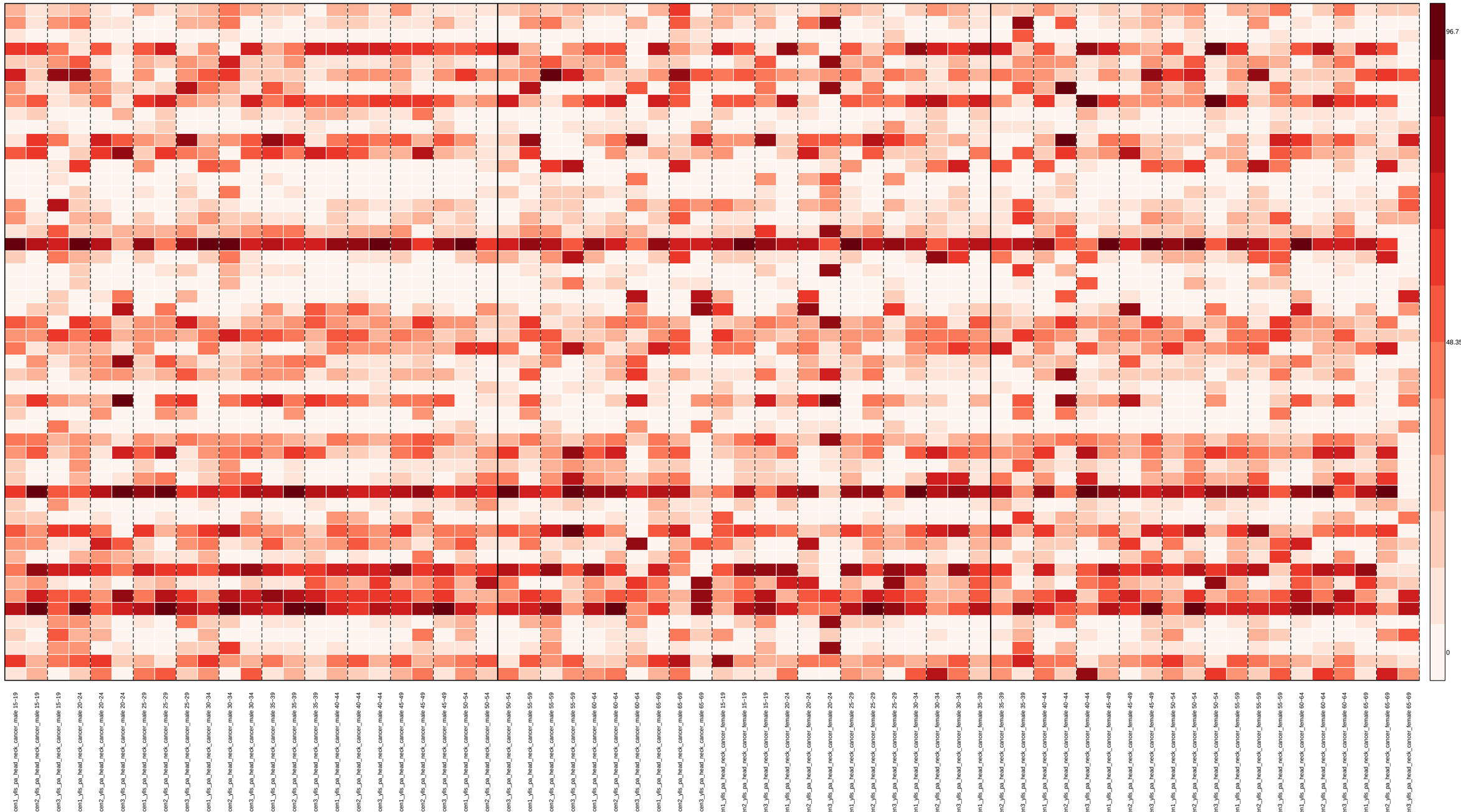
san_antonio − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



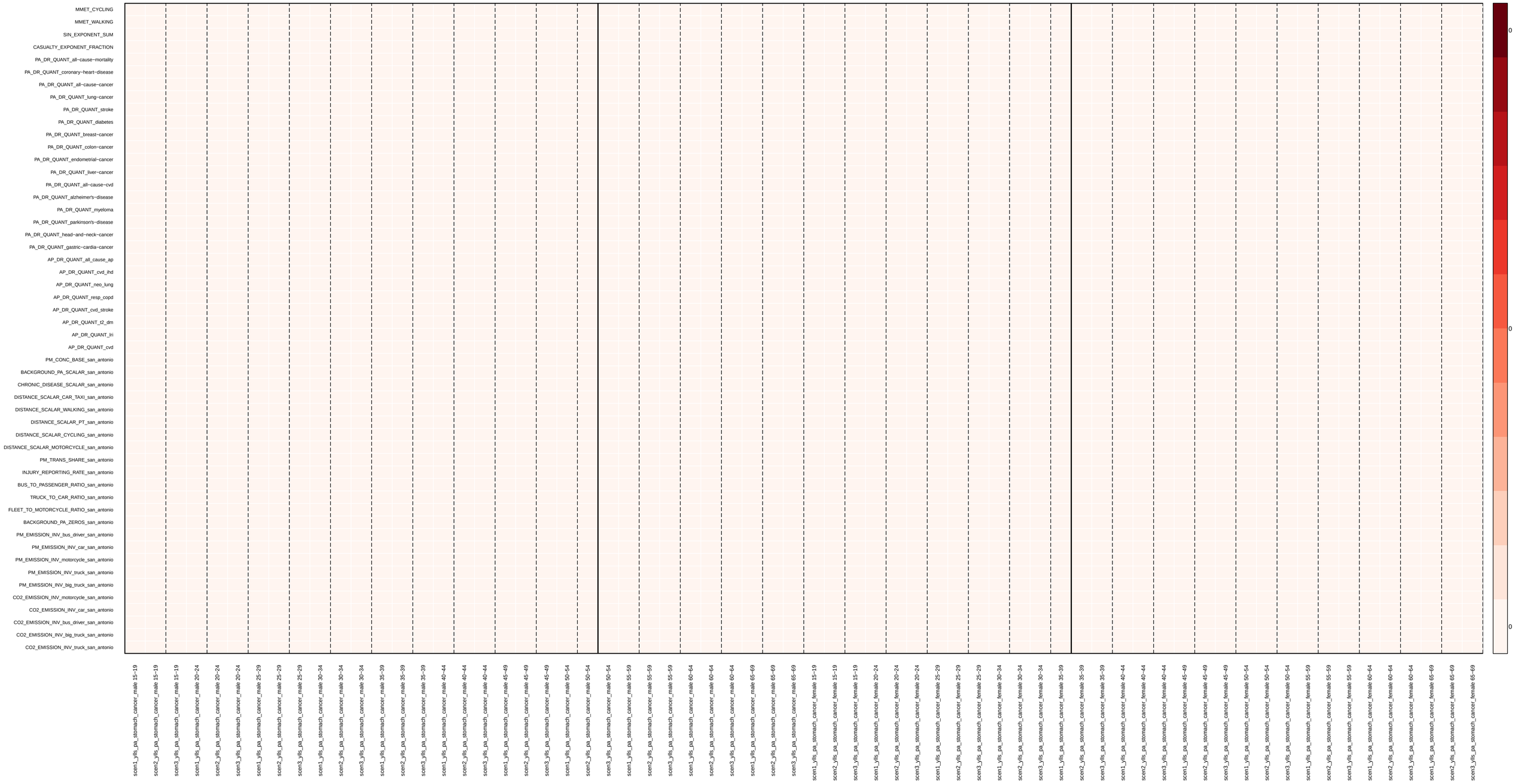
san_antonio − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



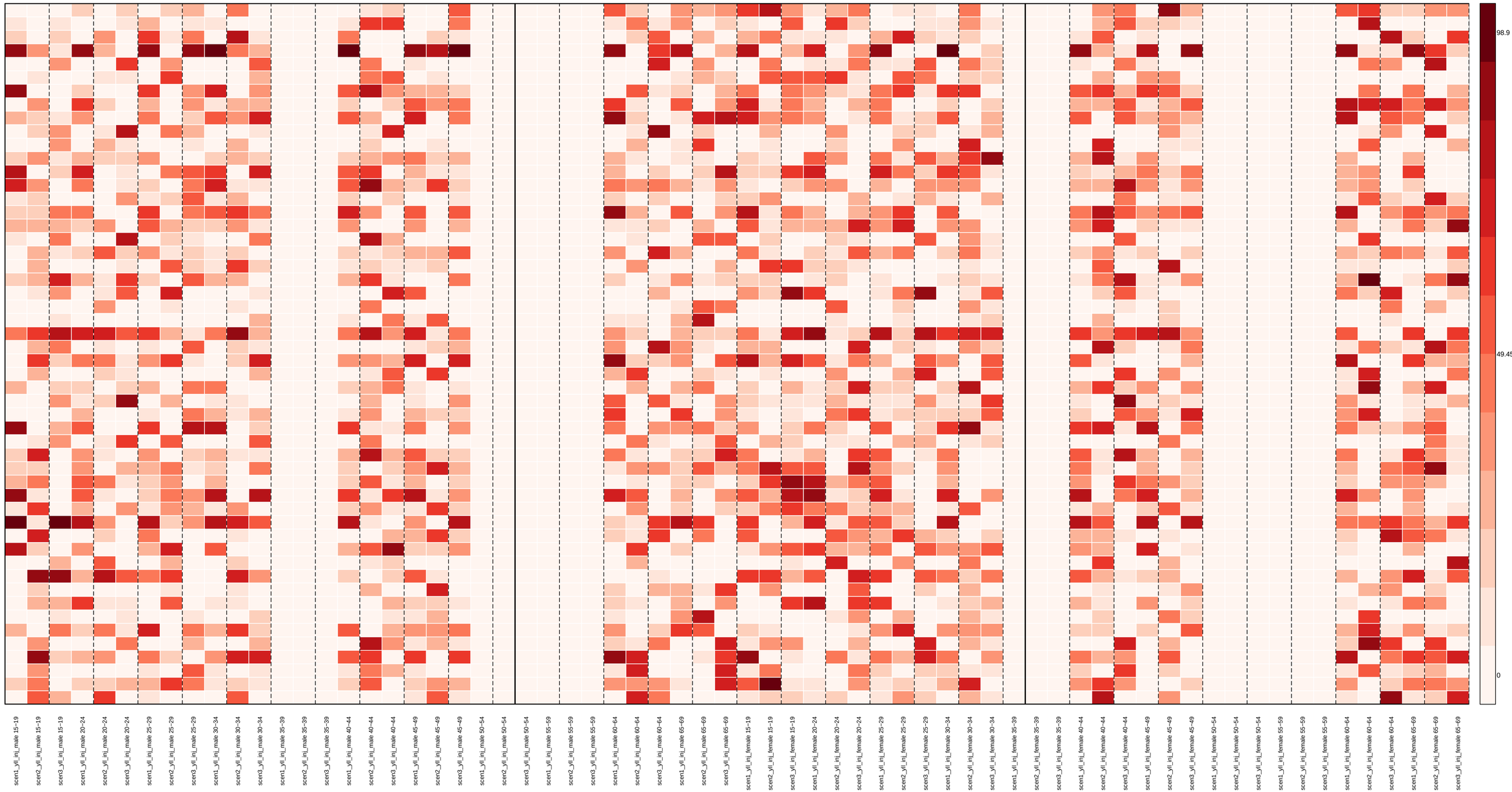
san_antonio − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



san_antonio − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

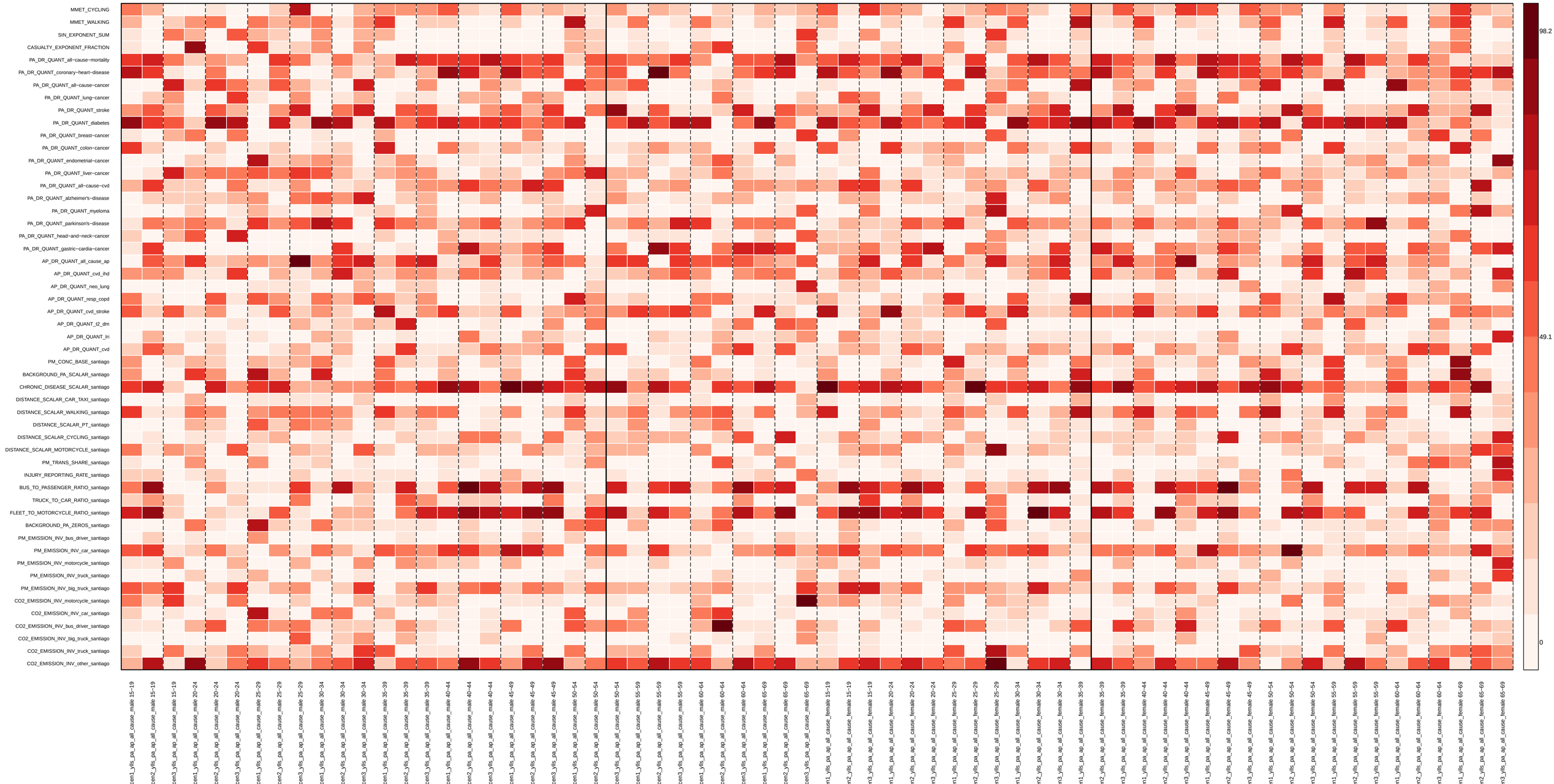


san_antonio − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

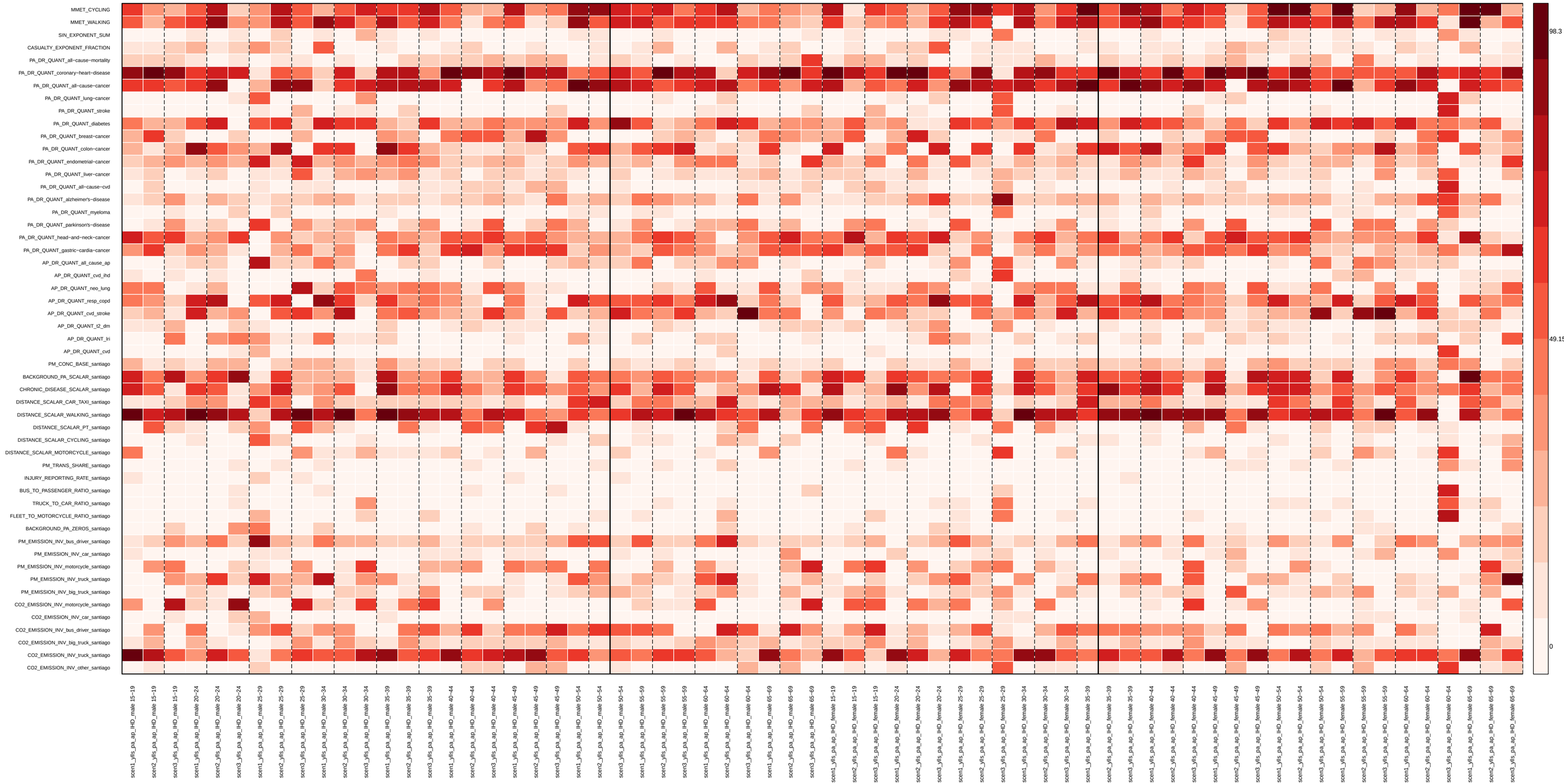


san_antonio − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

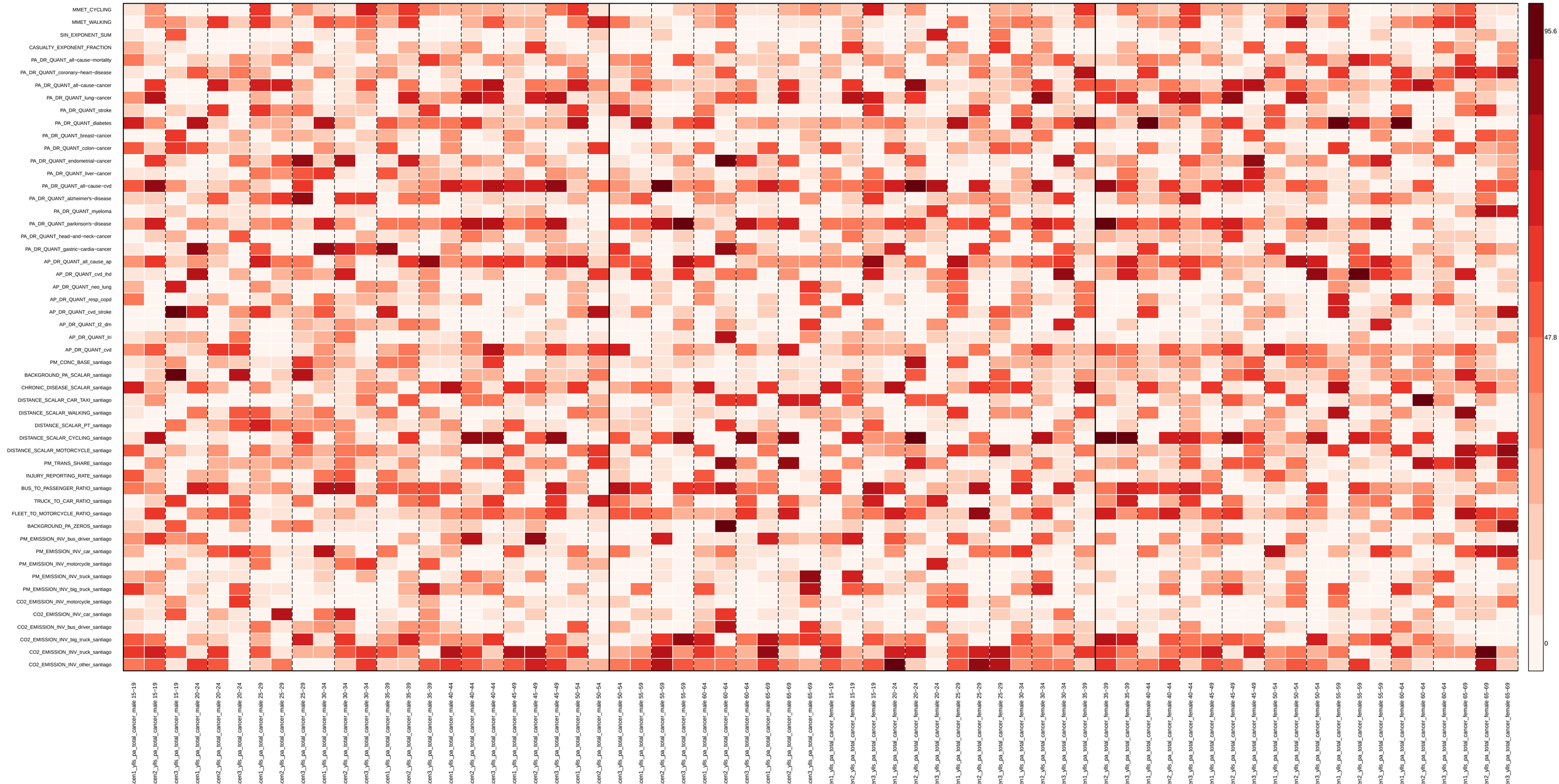
santiago − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



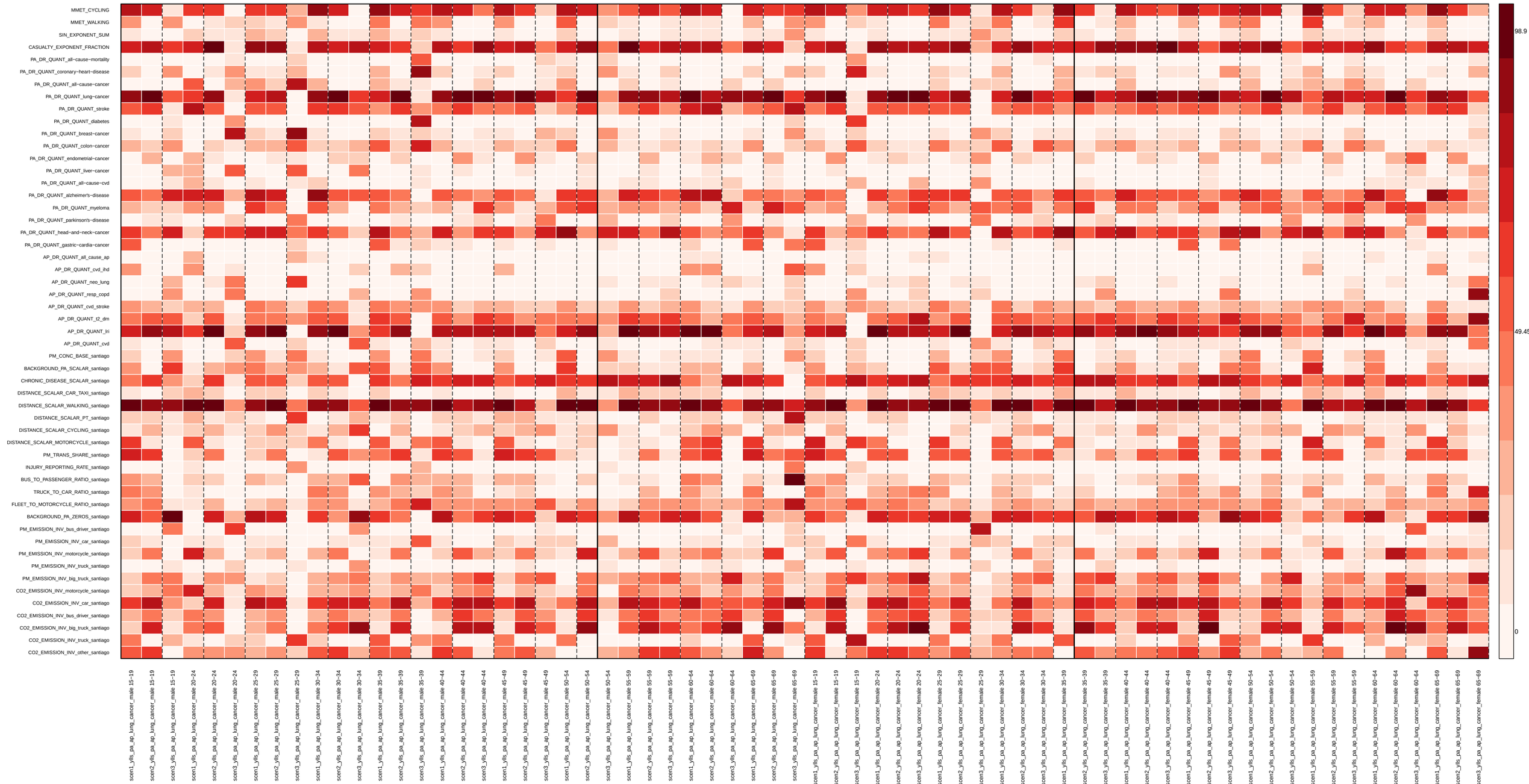
santiago − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



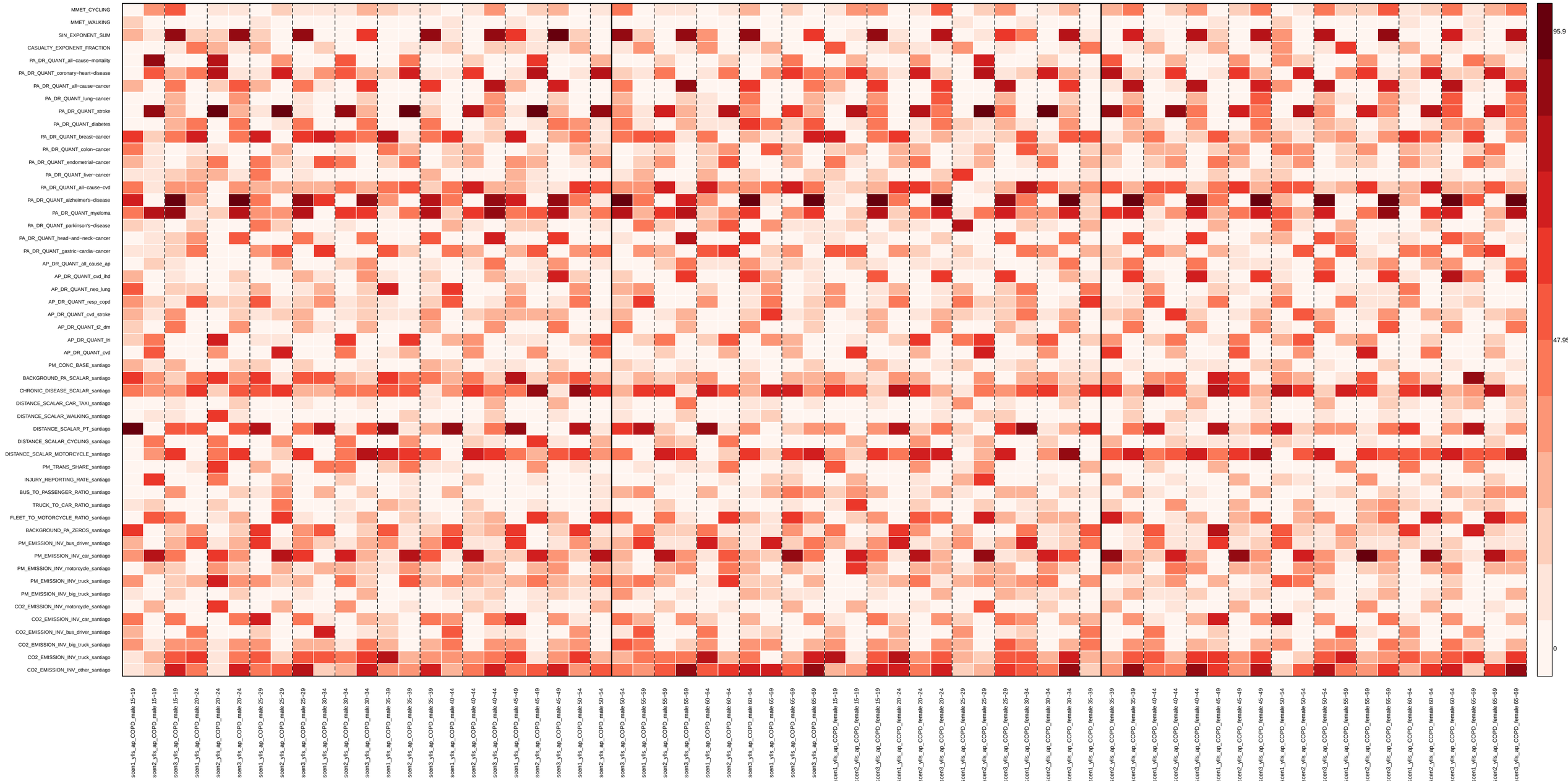
santiago − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



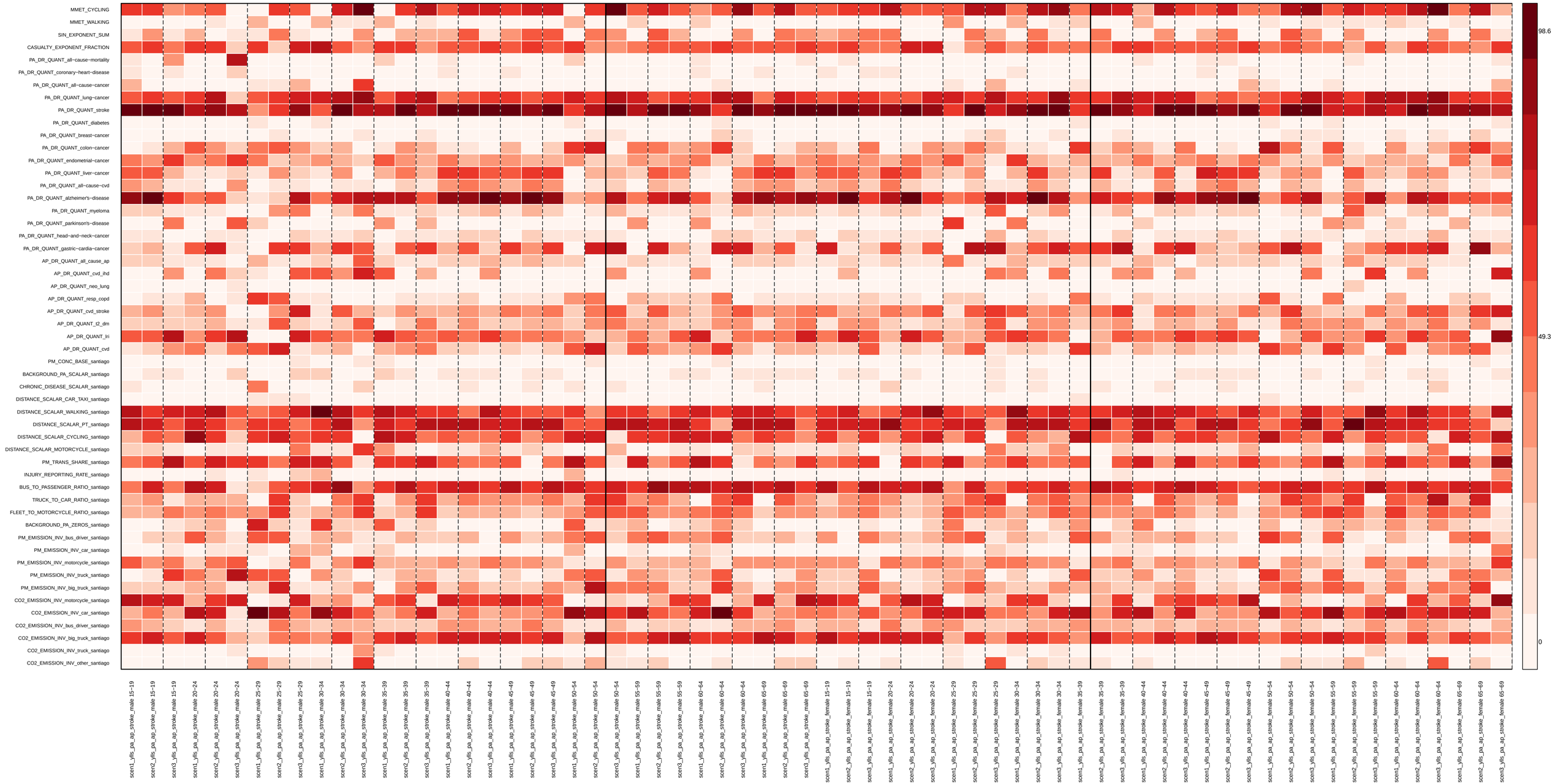
santiago − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



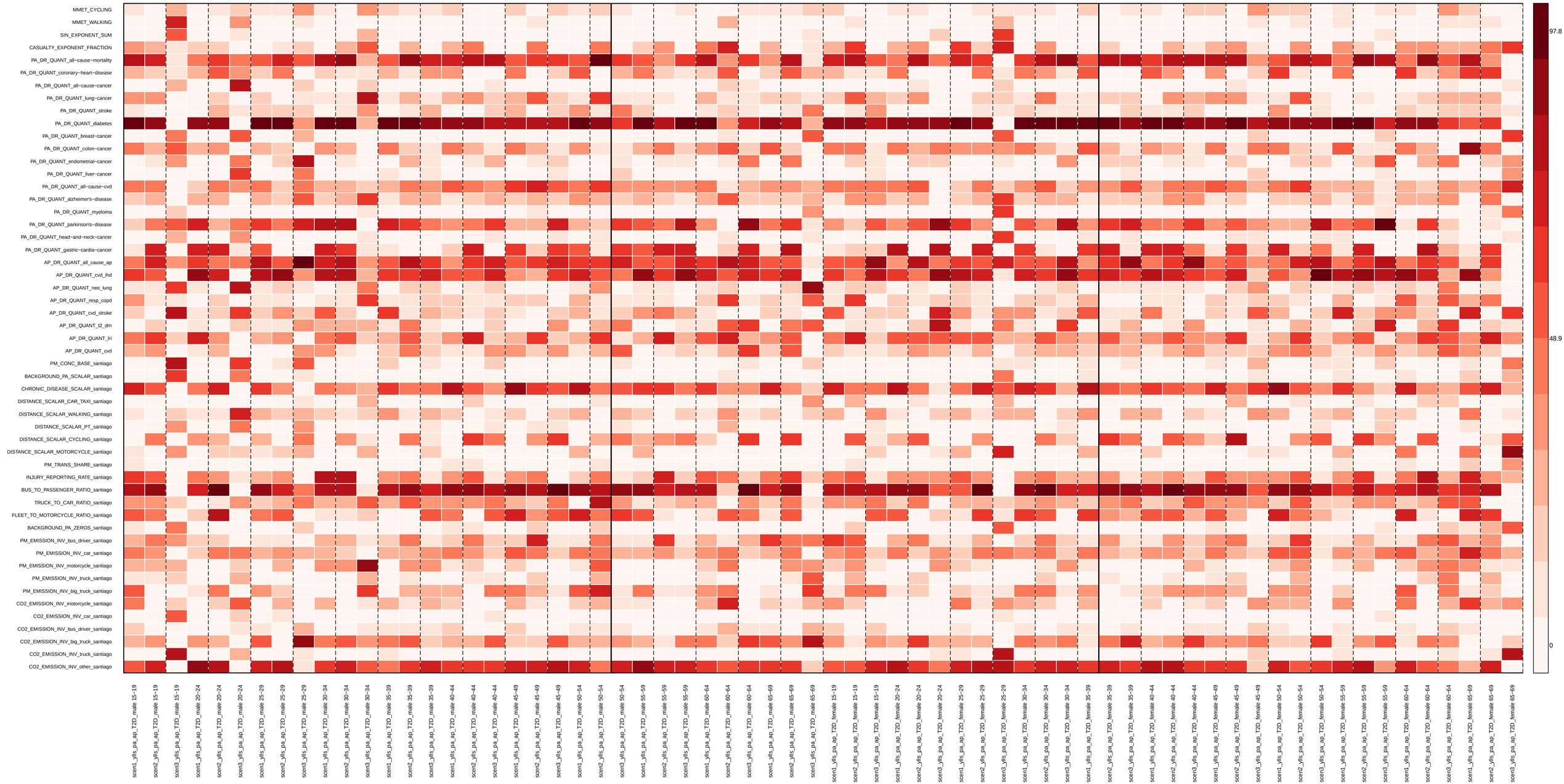
santiago − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



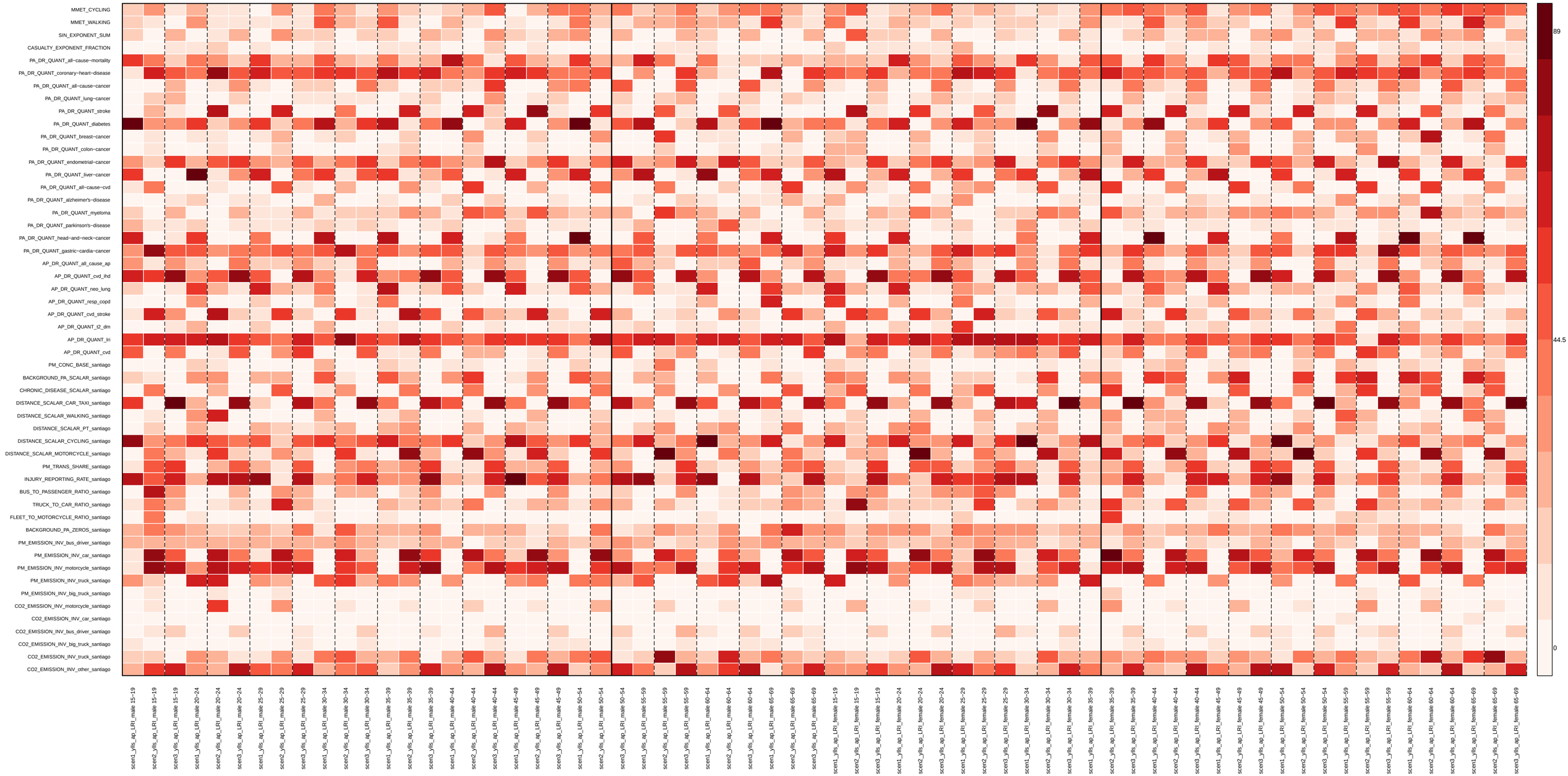
santiago − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



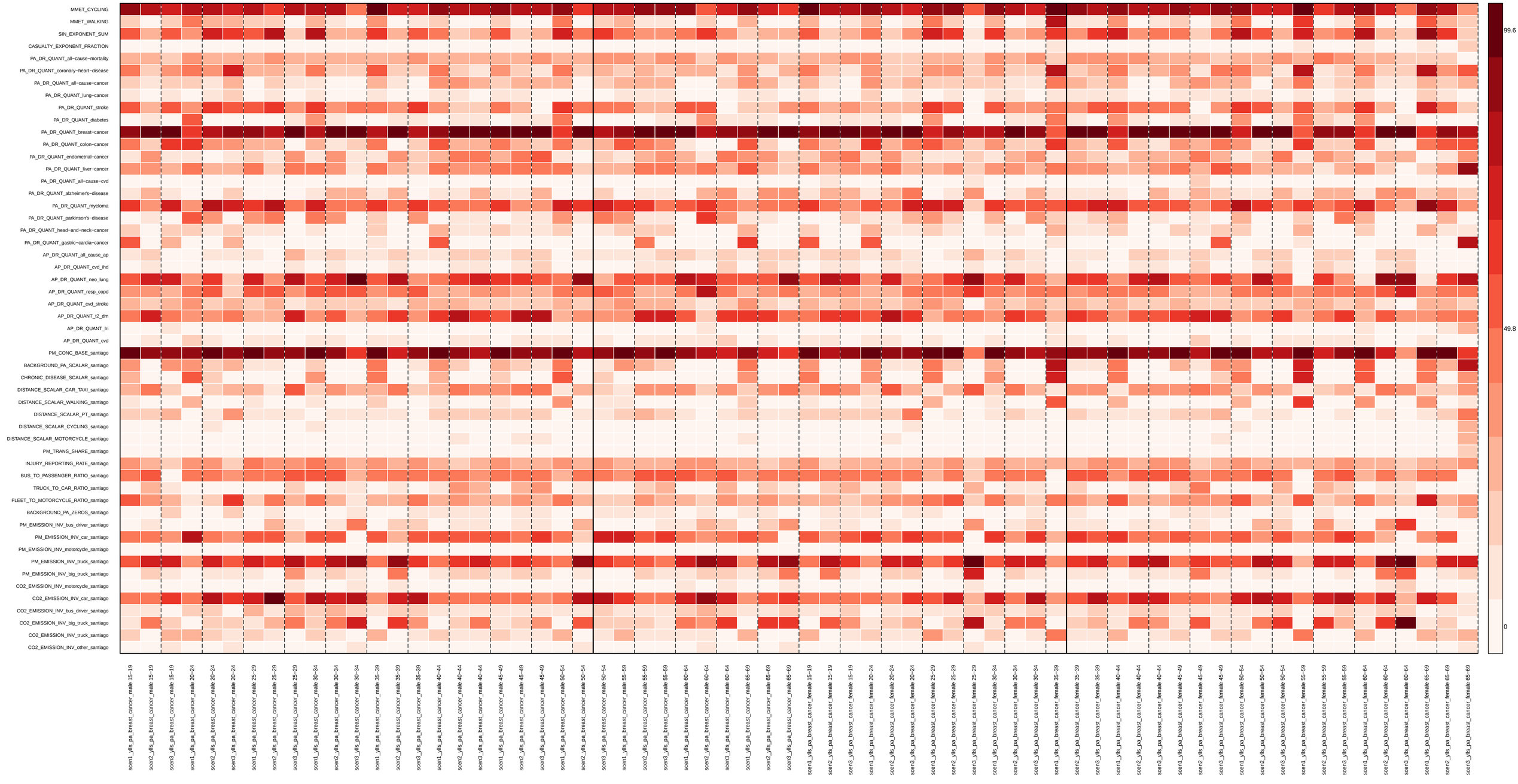
santiago − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



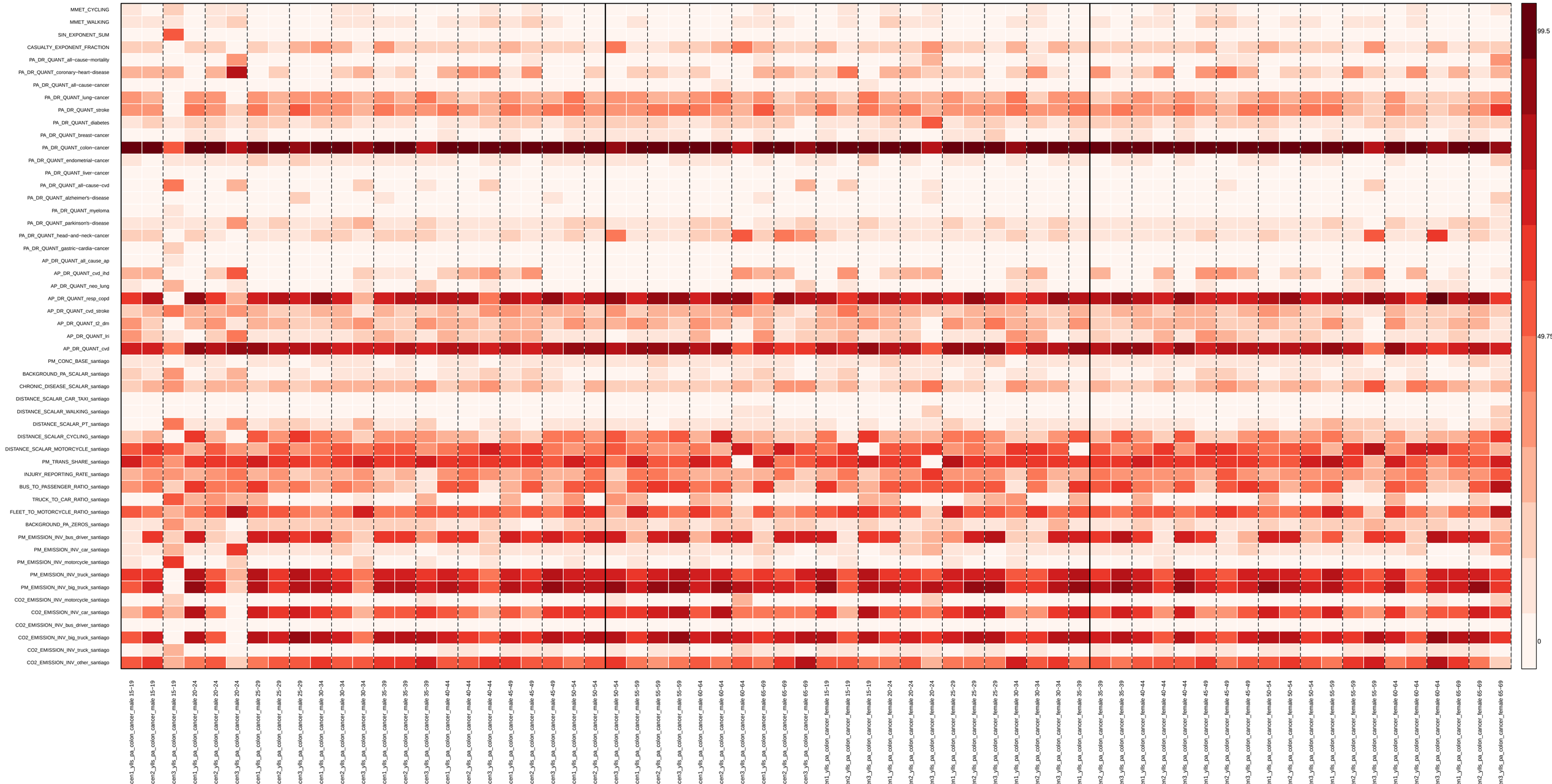
santiago − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



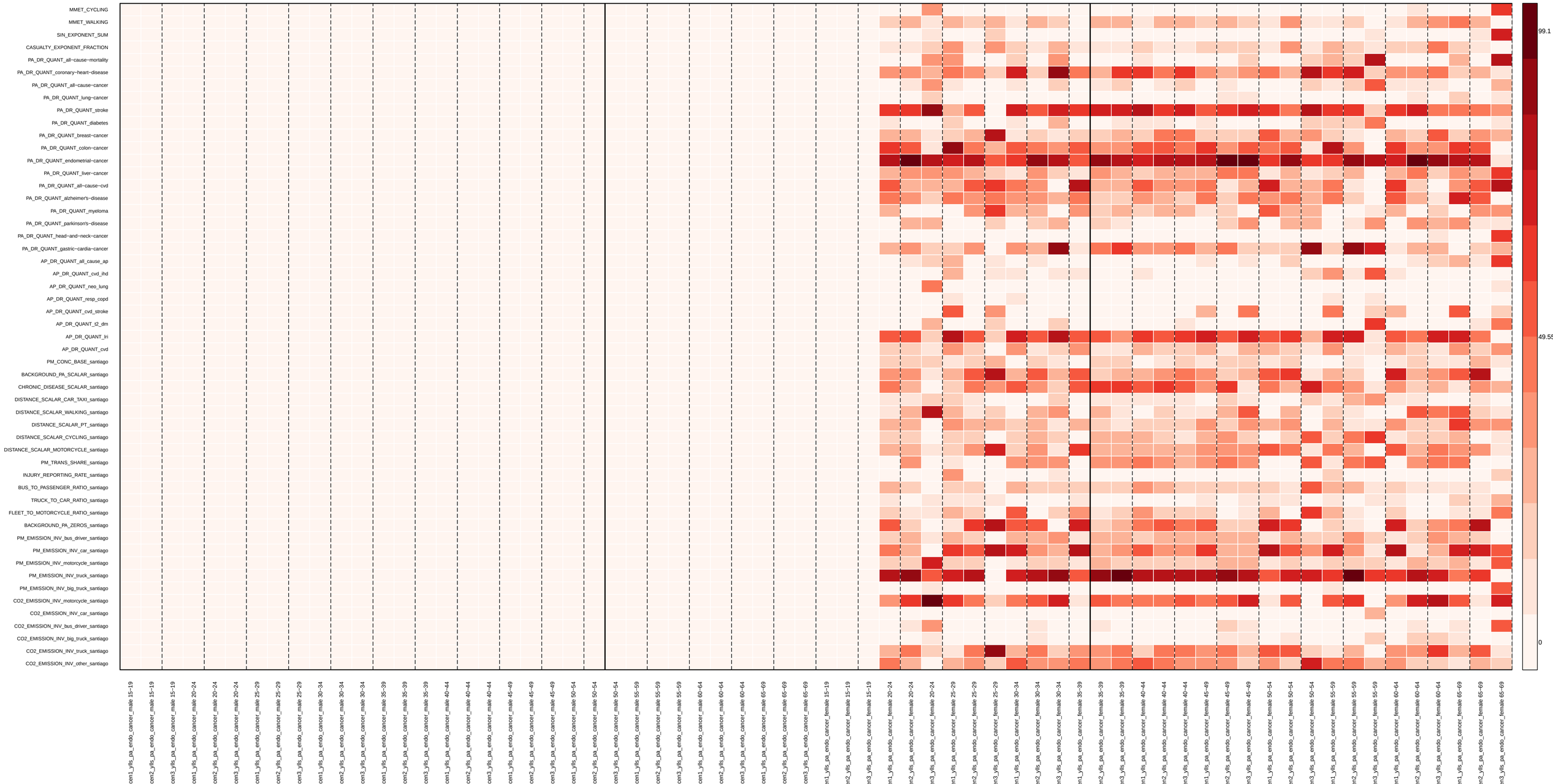
santiago − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



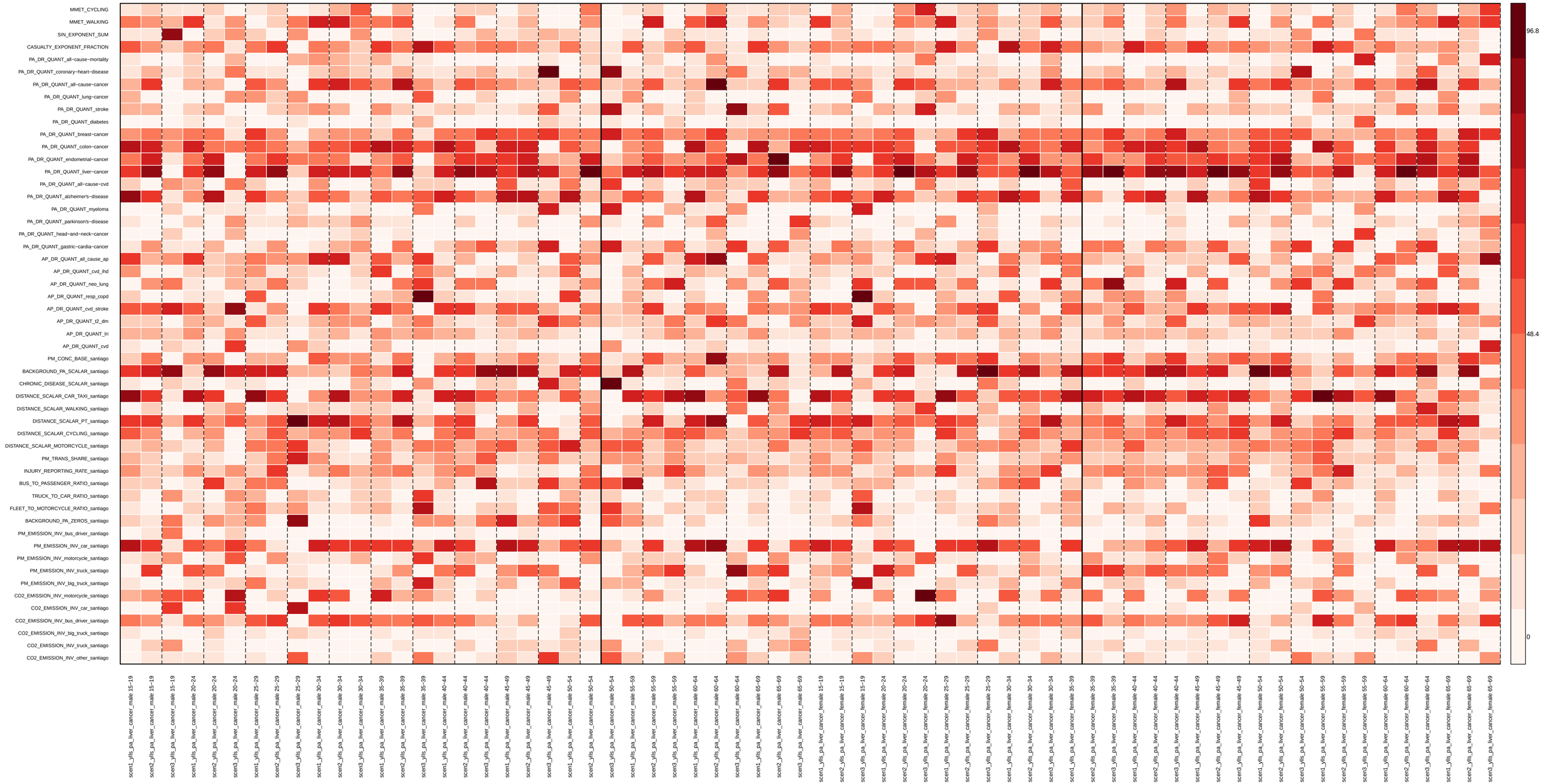
santiago − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



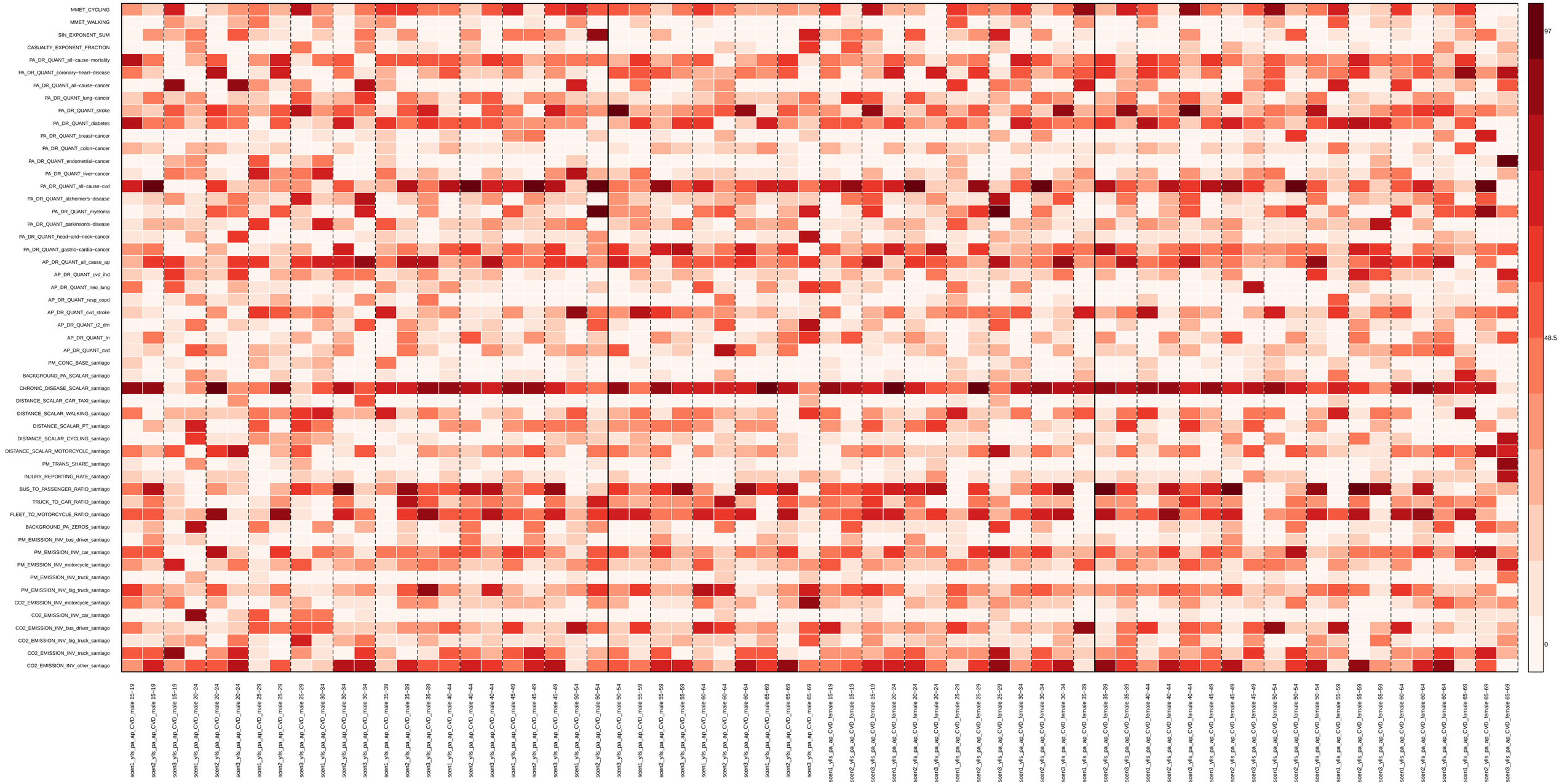
santiago − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



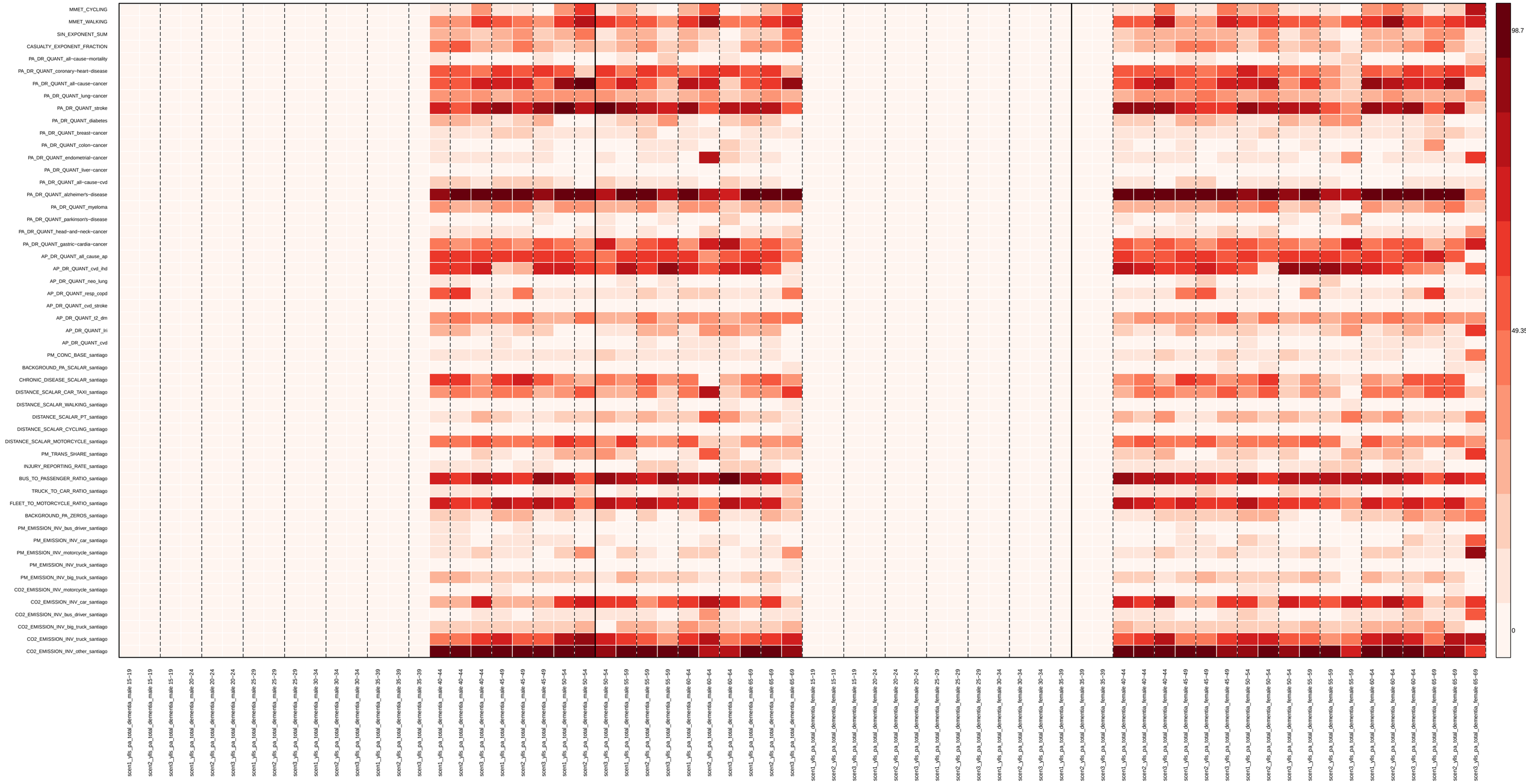
santiago − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



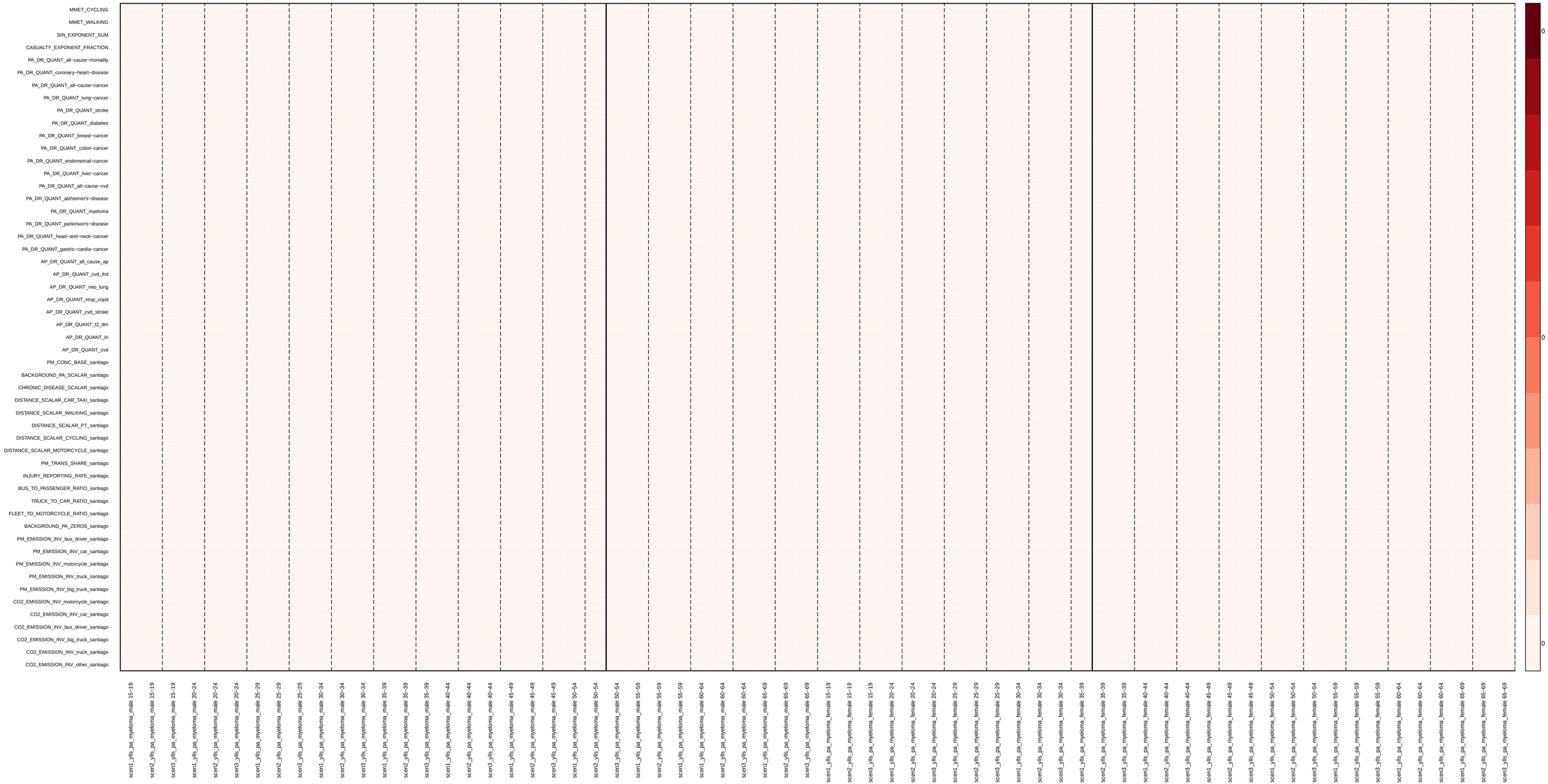
santiago − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



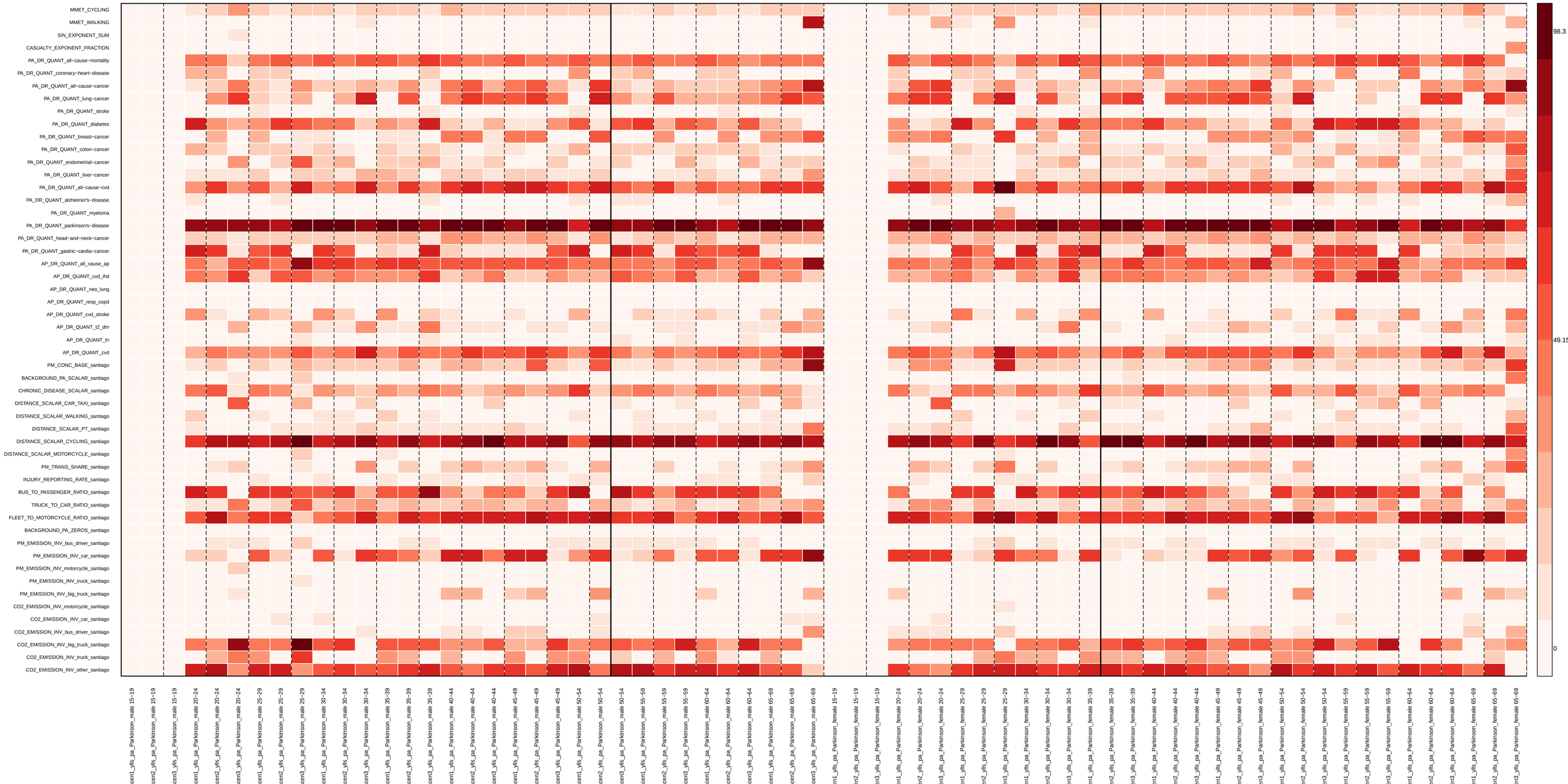
santiago − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



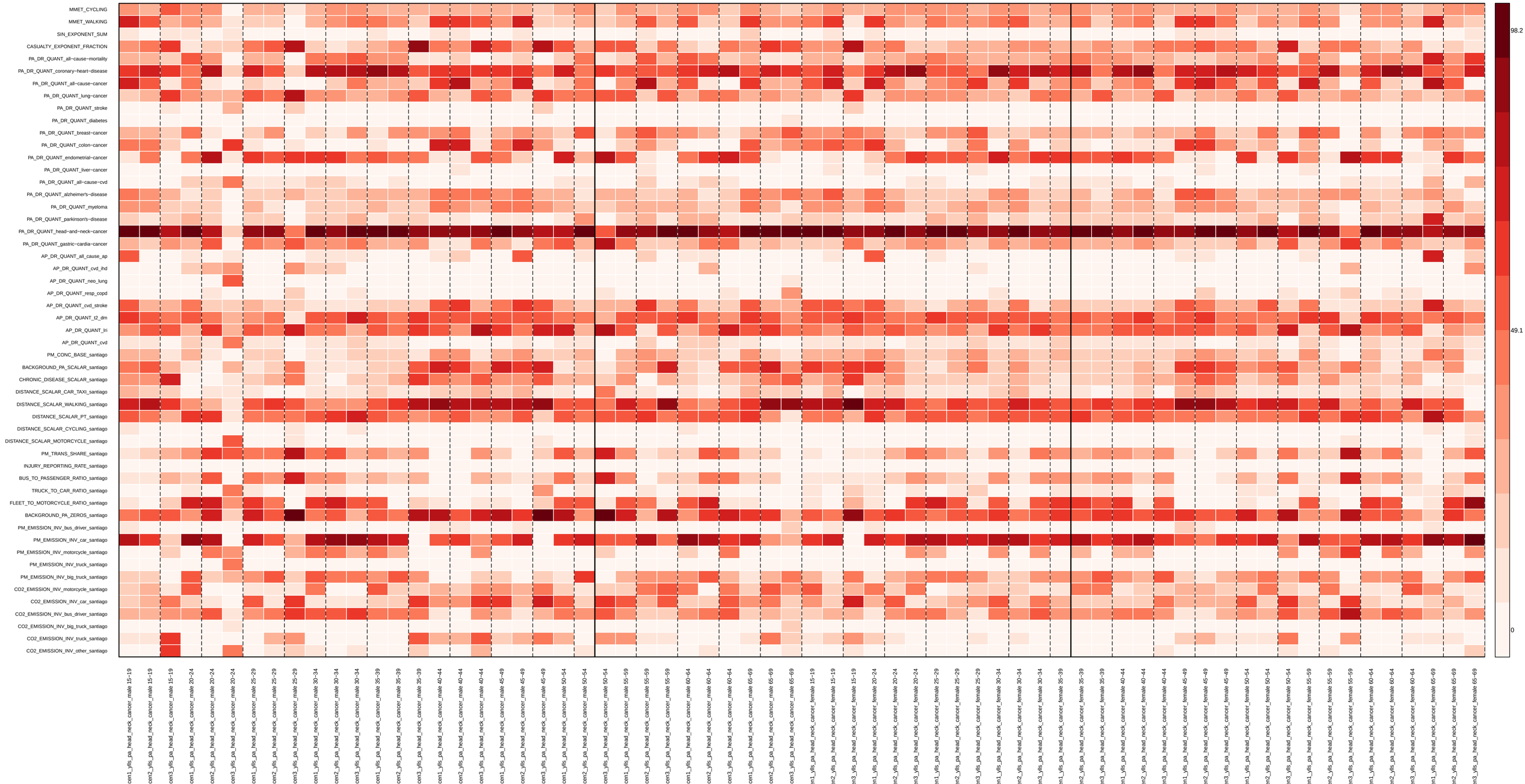
santiago − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



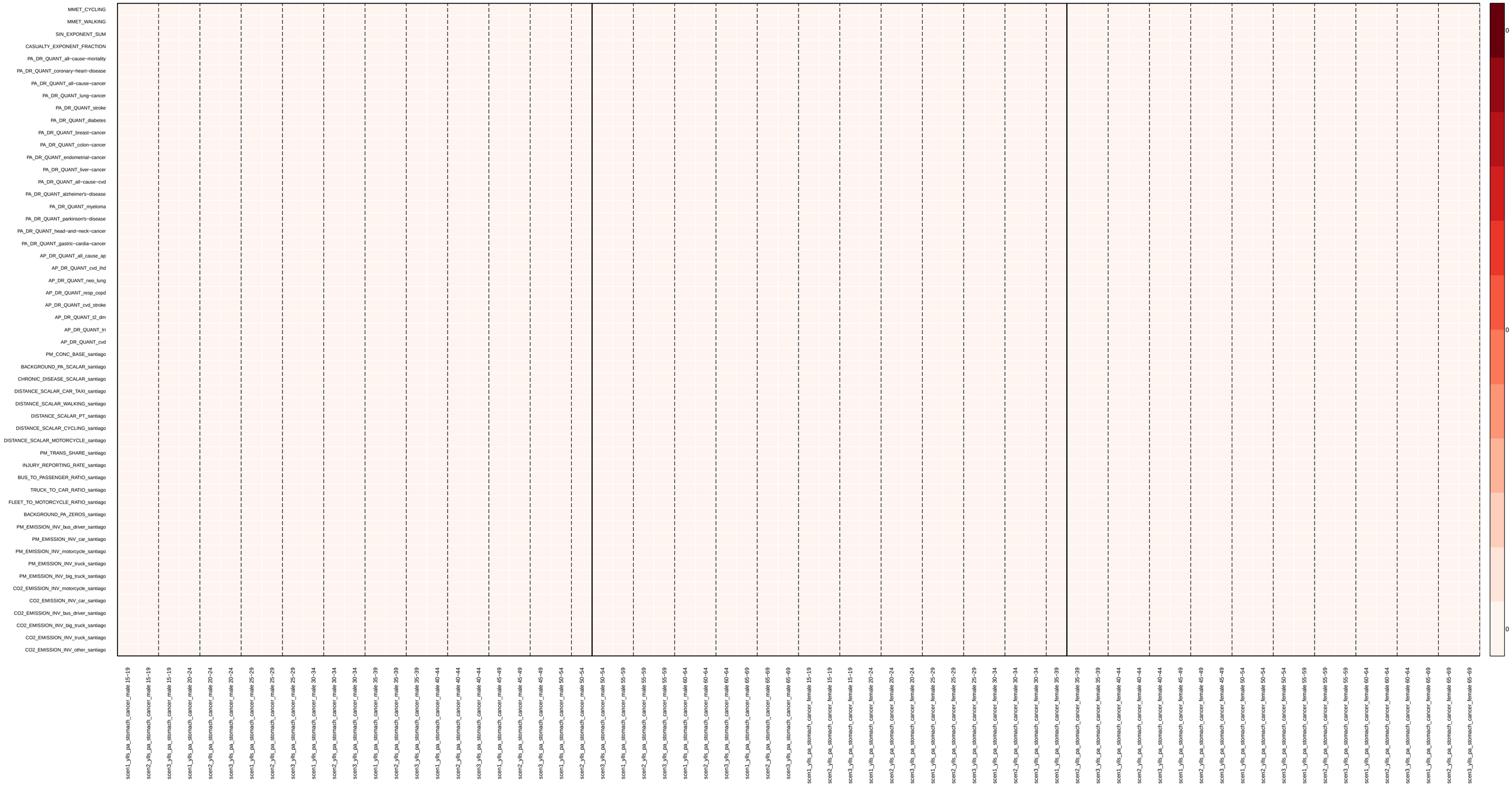
santiago − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



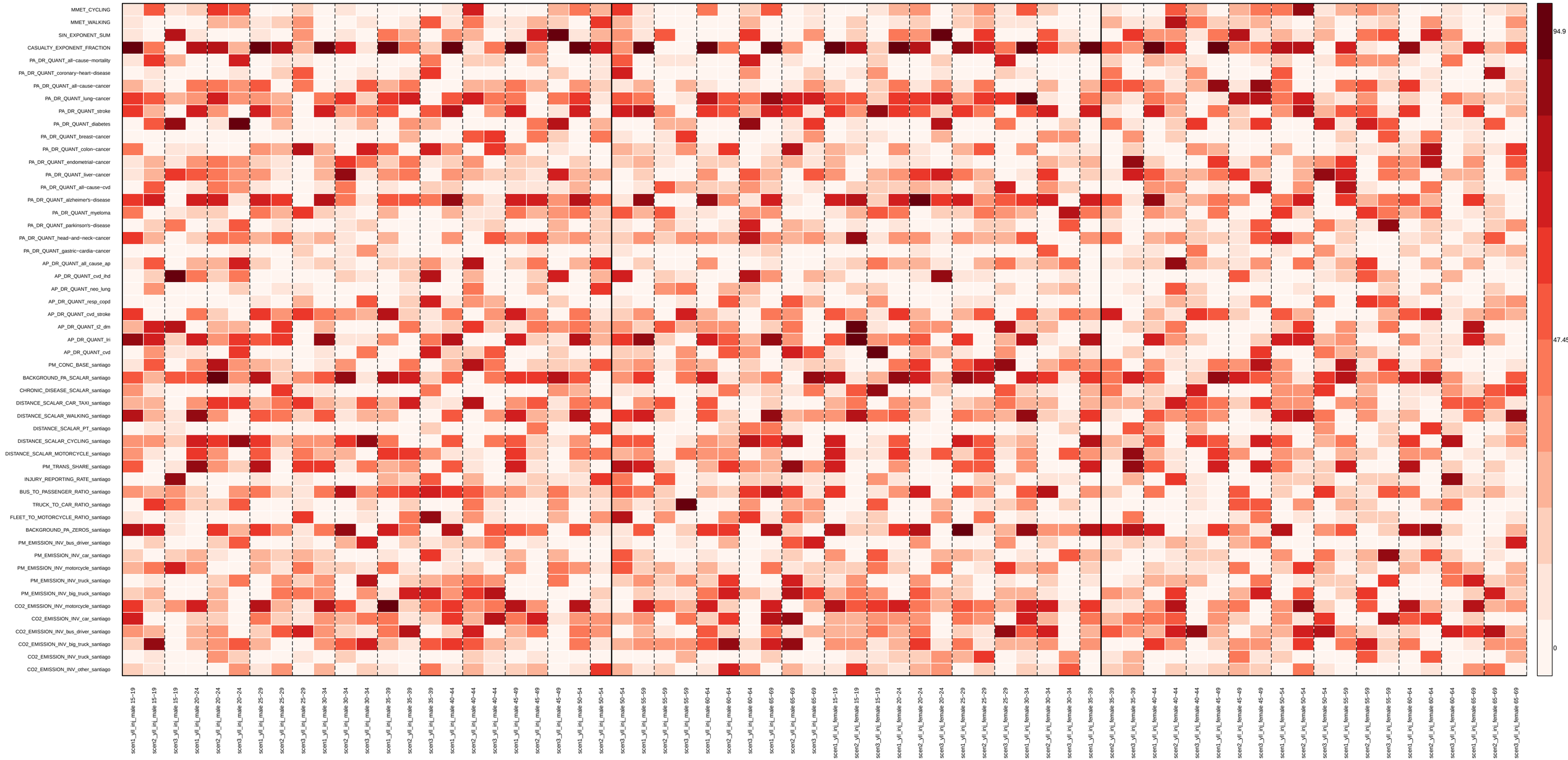
santiago − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



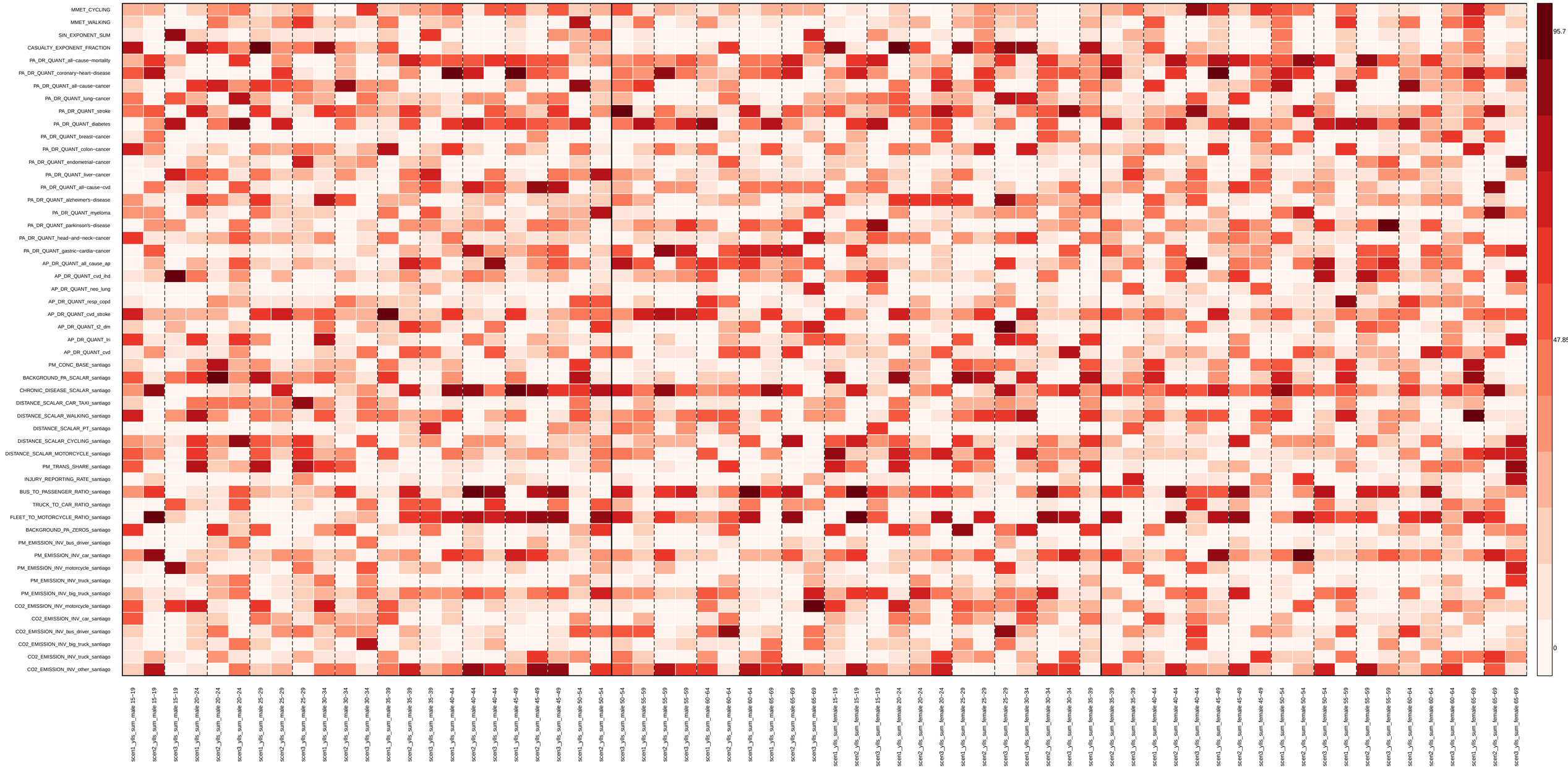
santiago − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



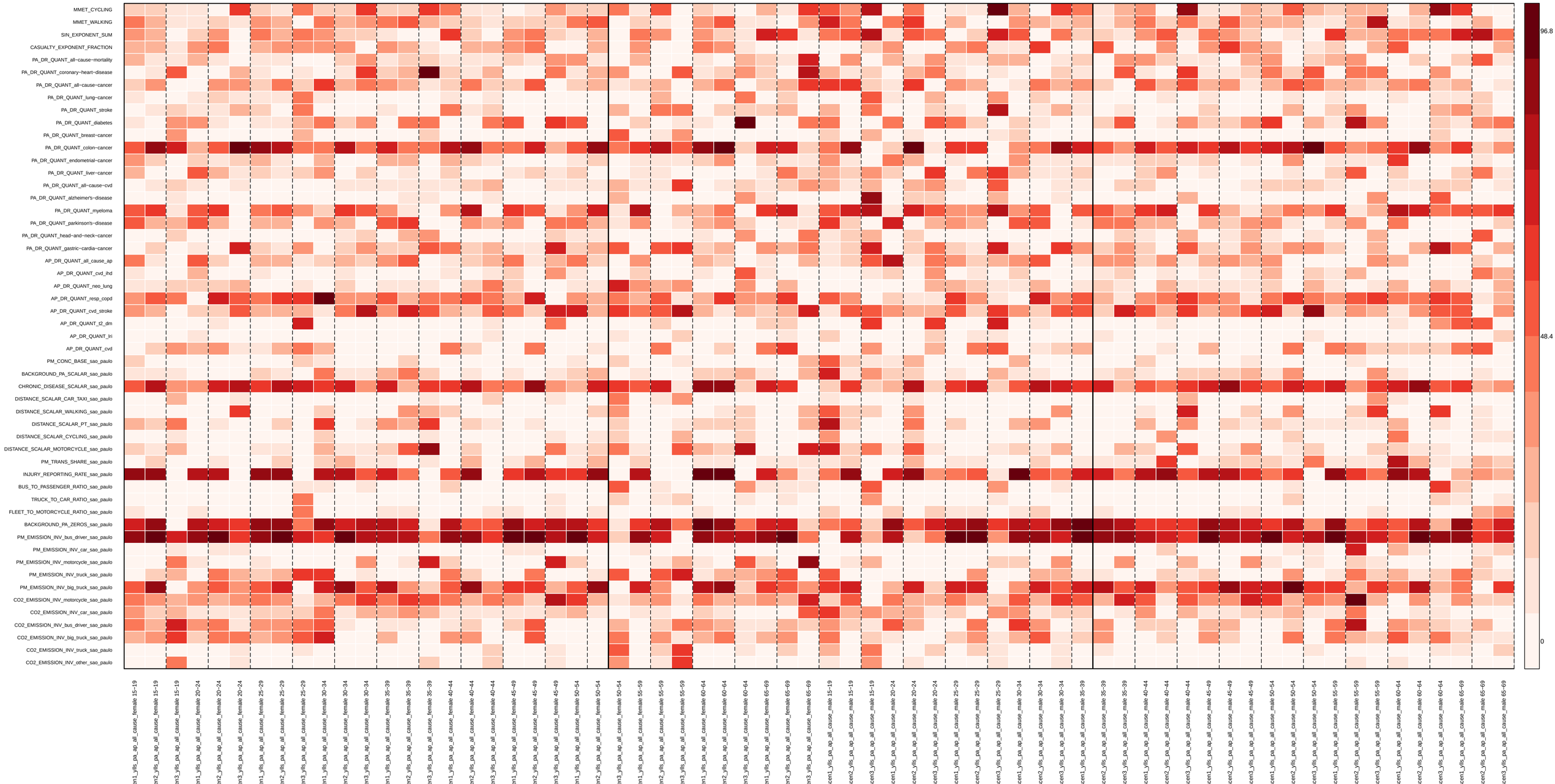
santiago − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



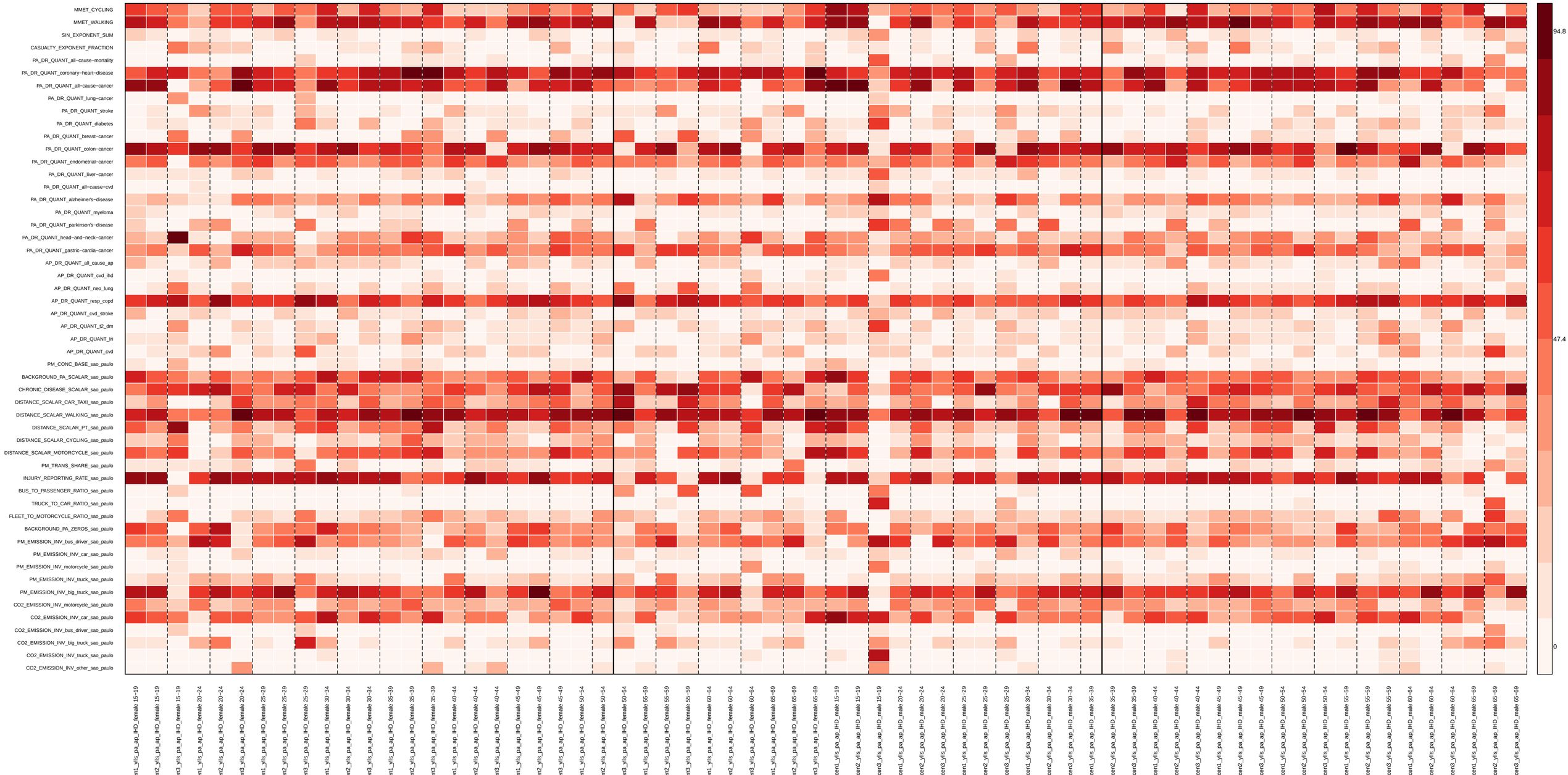
santiago − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



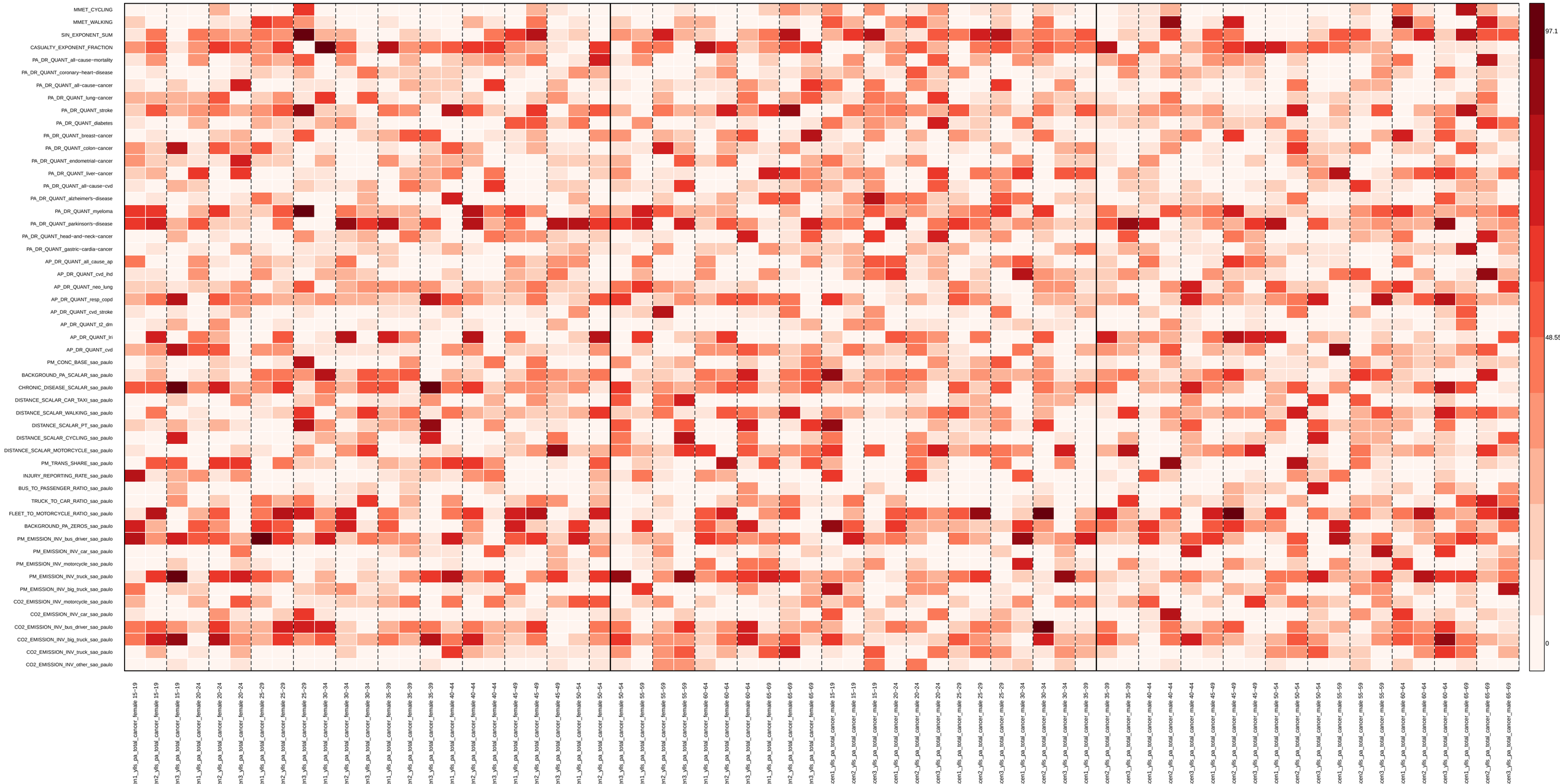
sao_paulo − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



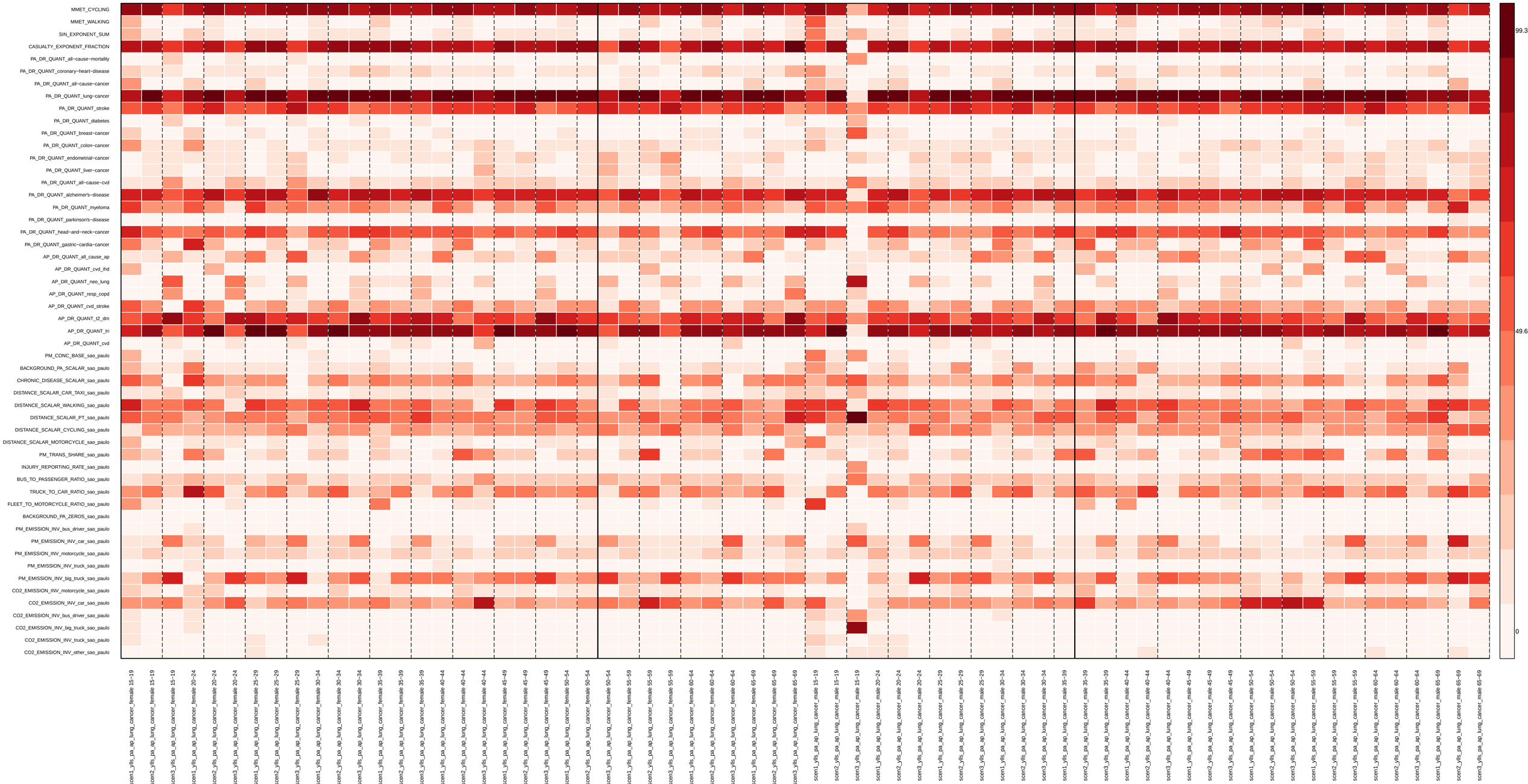
sao_paulo − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



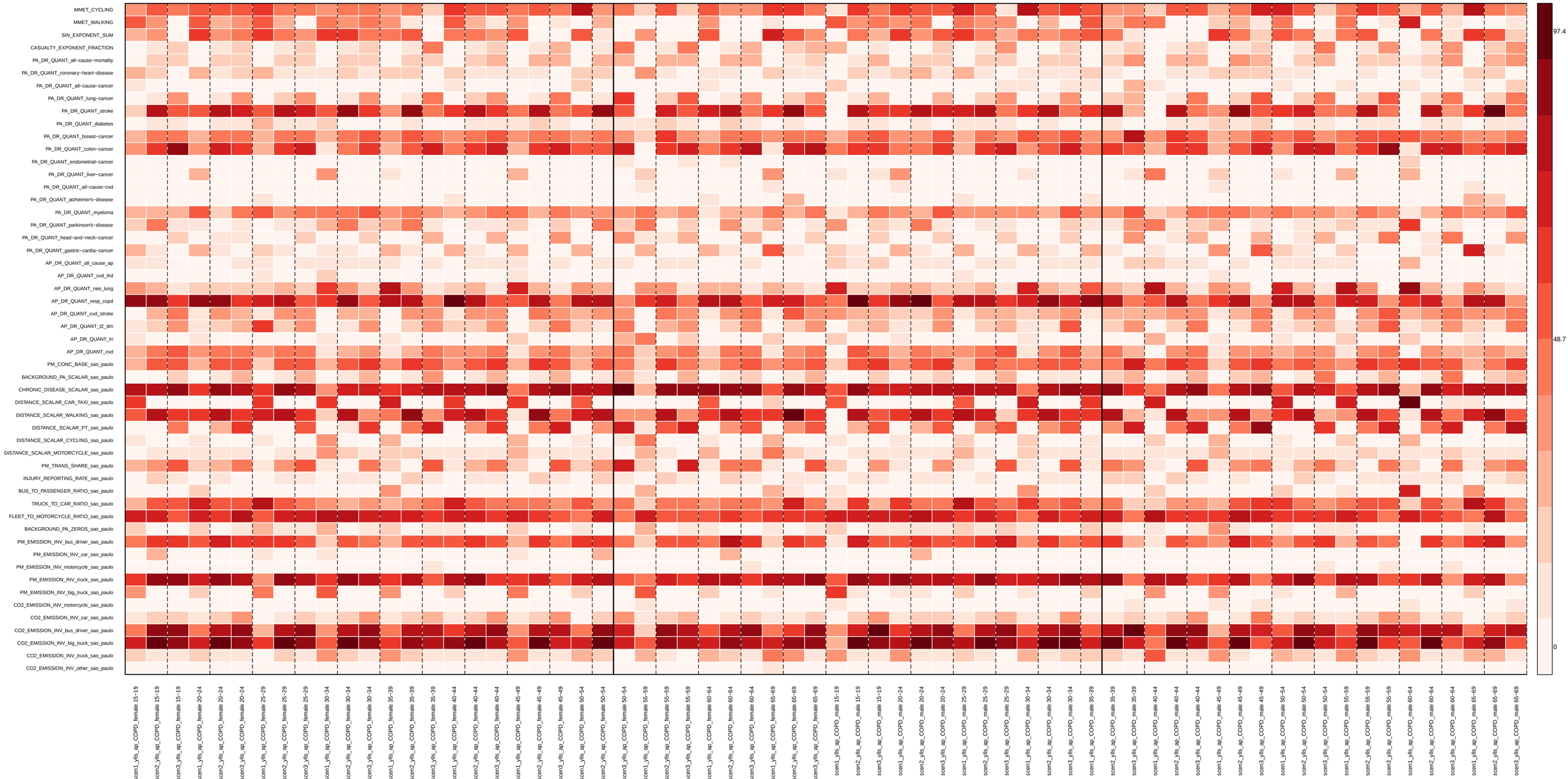
sao_paulo − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



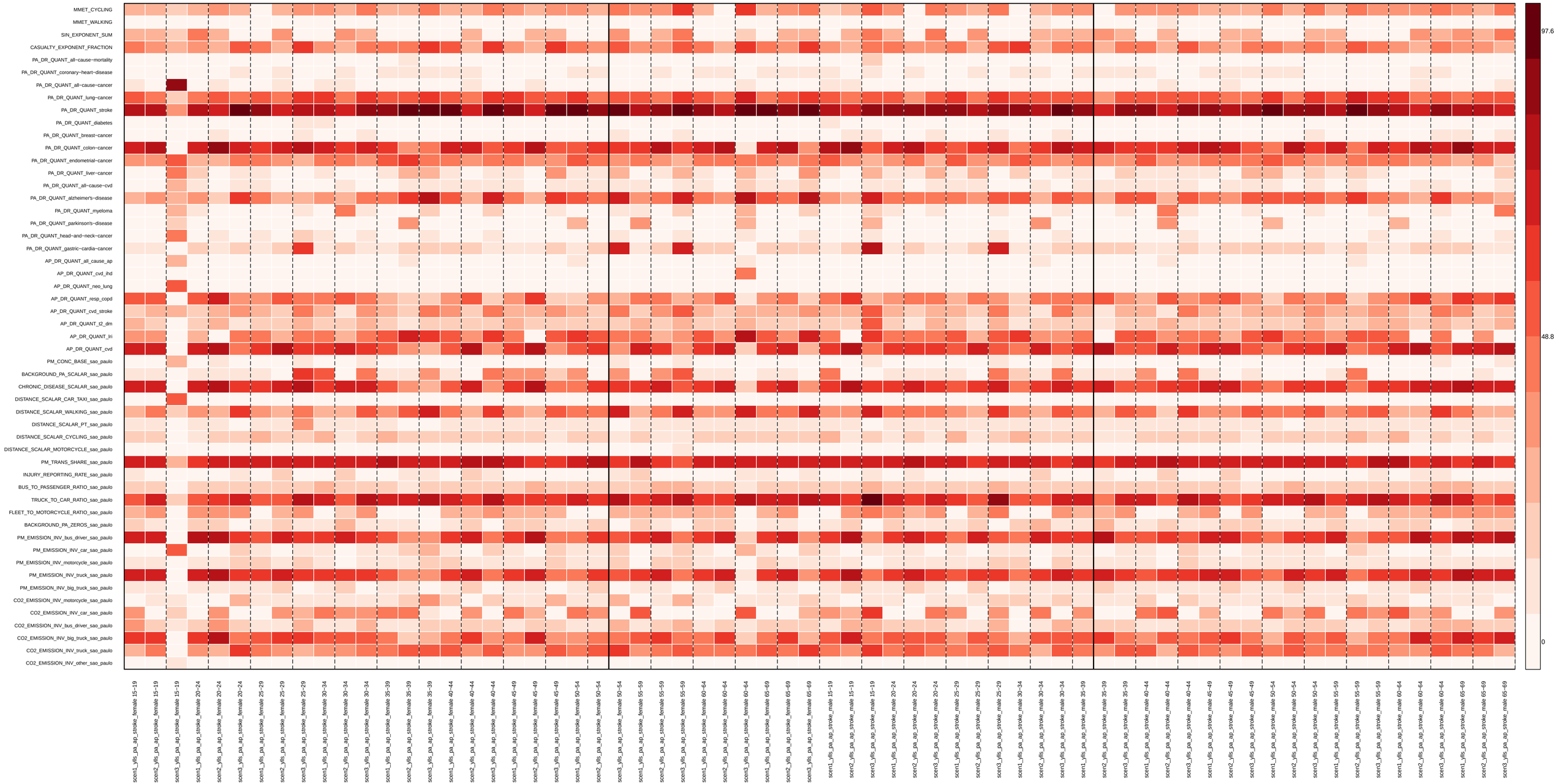
sao_paulo − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



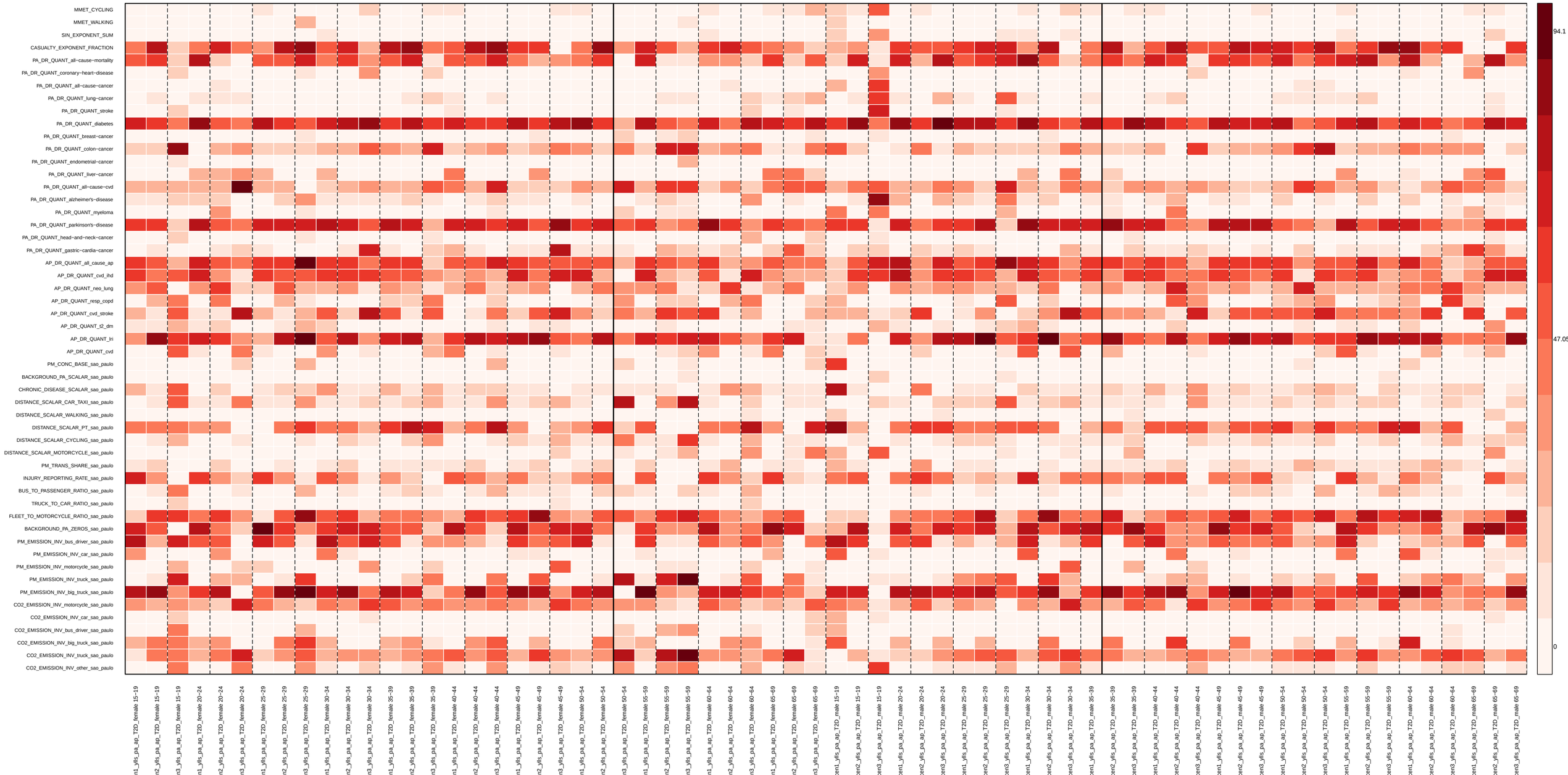
sao_paulo − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



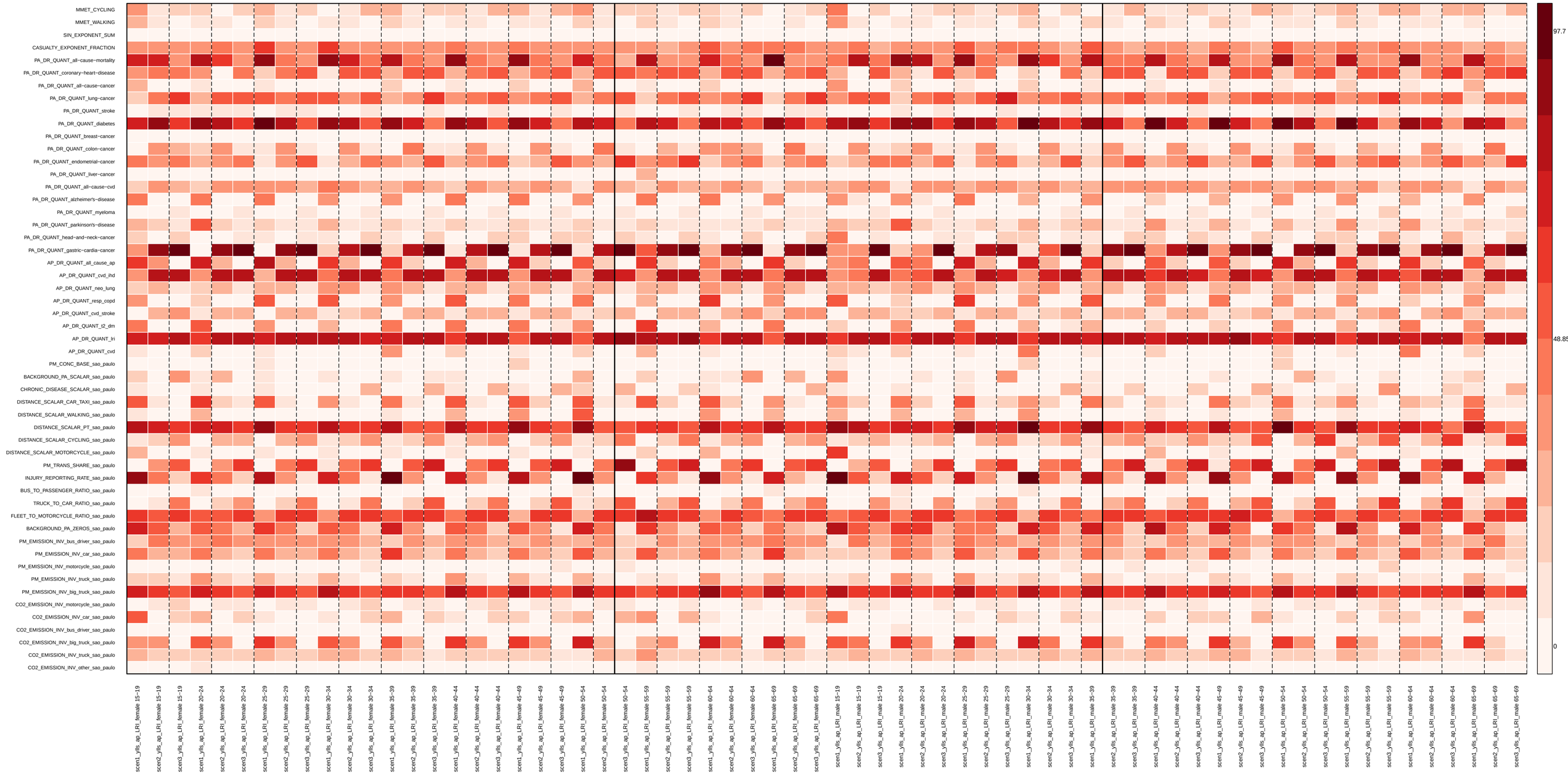
sao_paulo − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



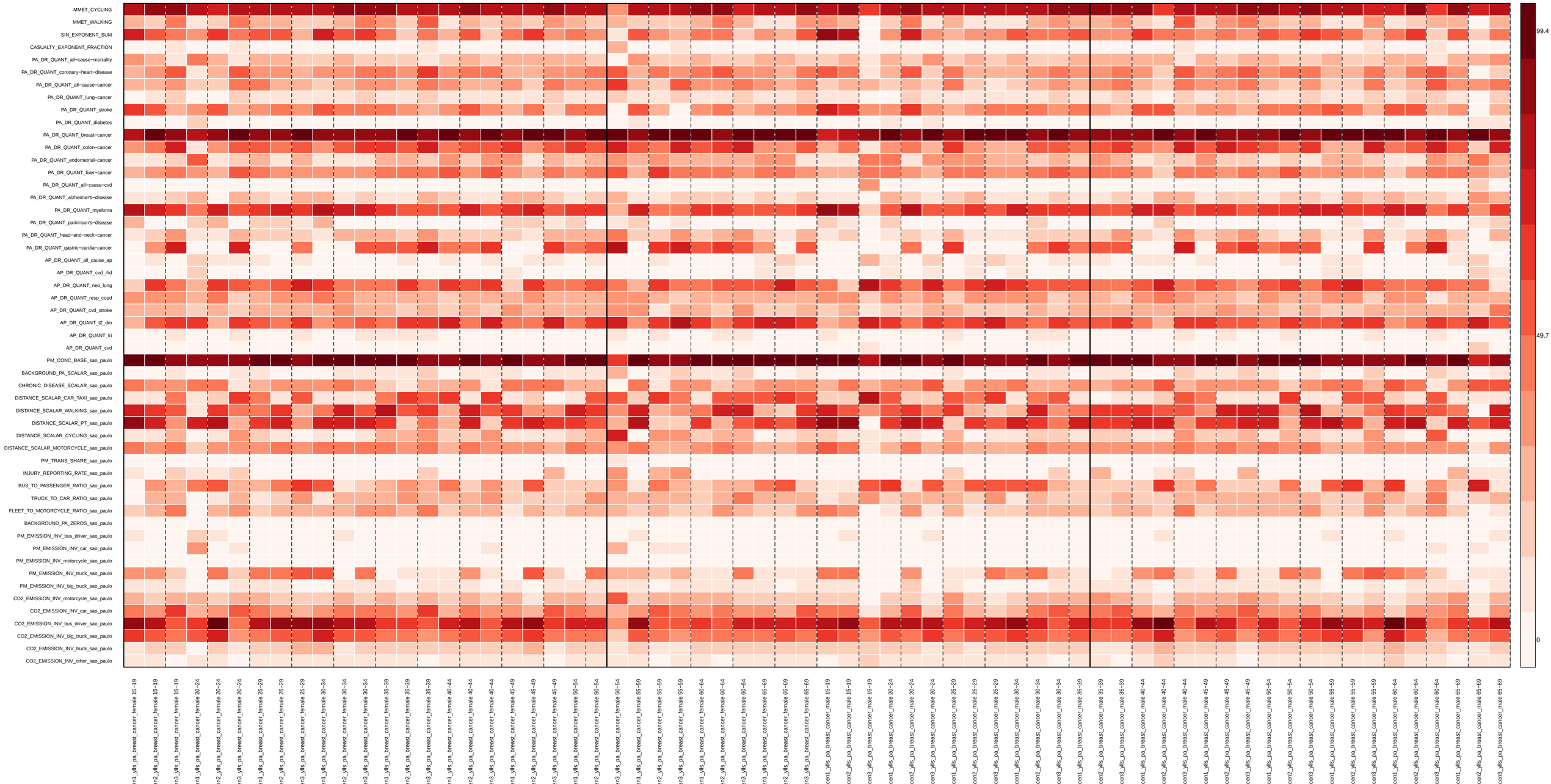
sao_paulo − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



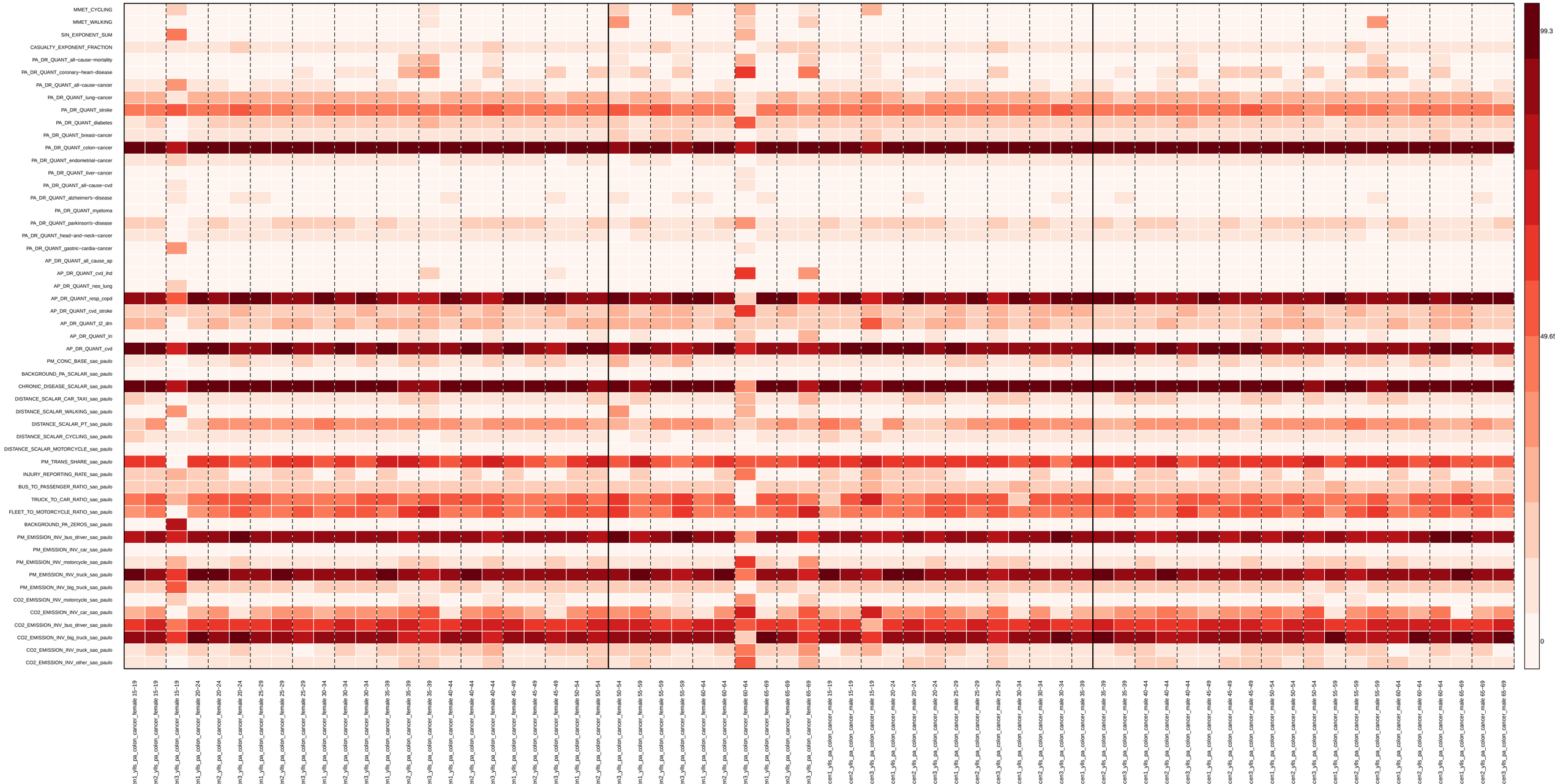
sao_paulo − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



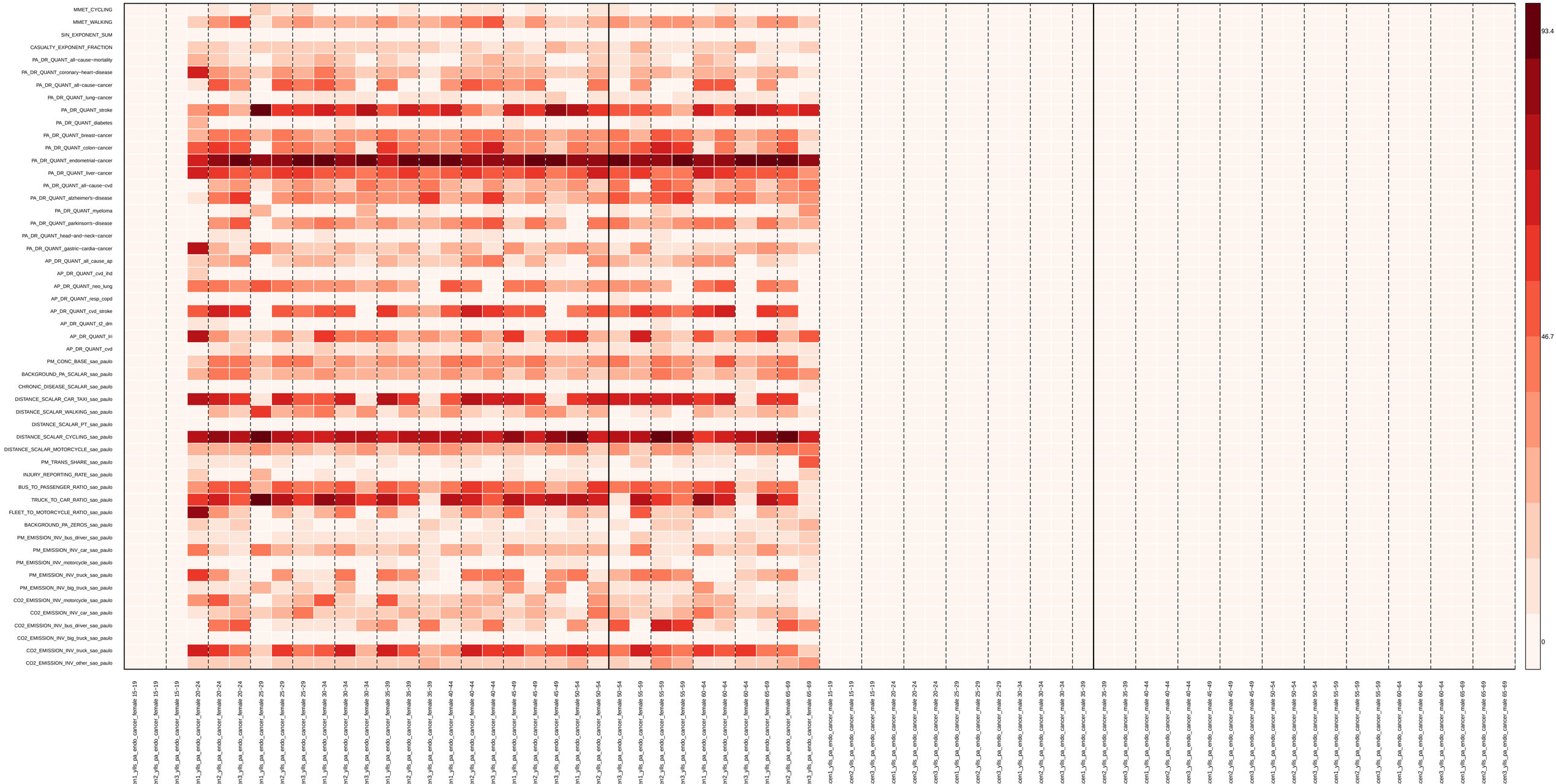
sao_paulo − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



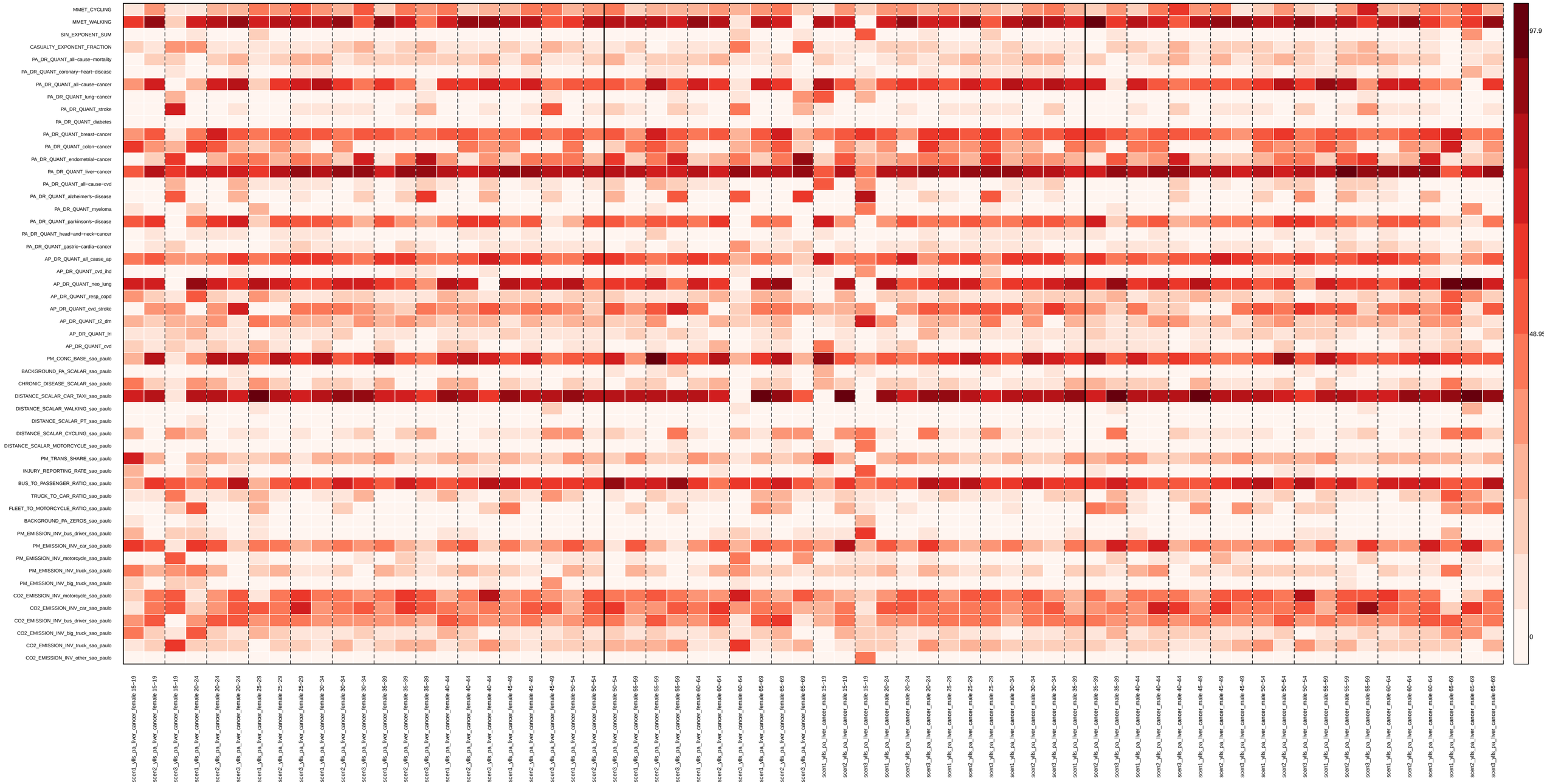
sao_paulo − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



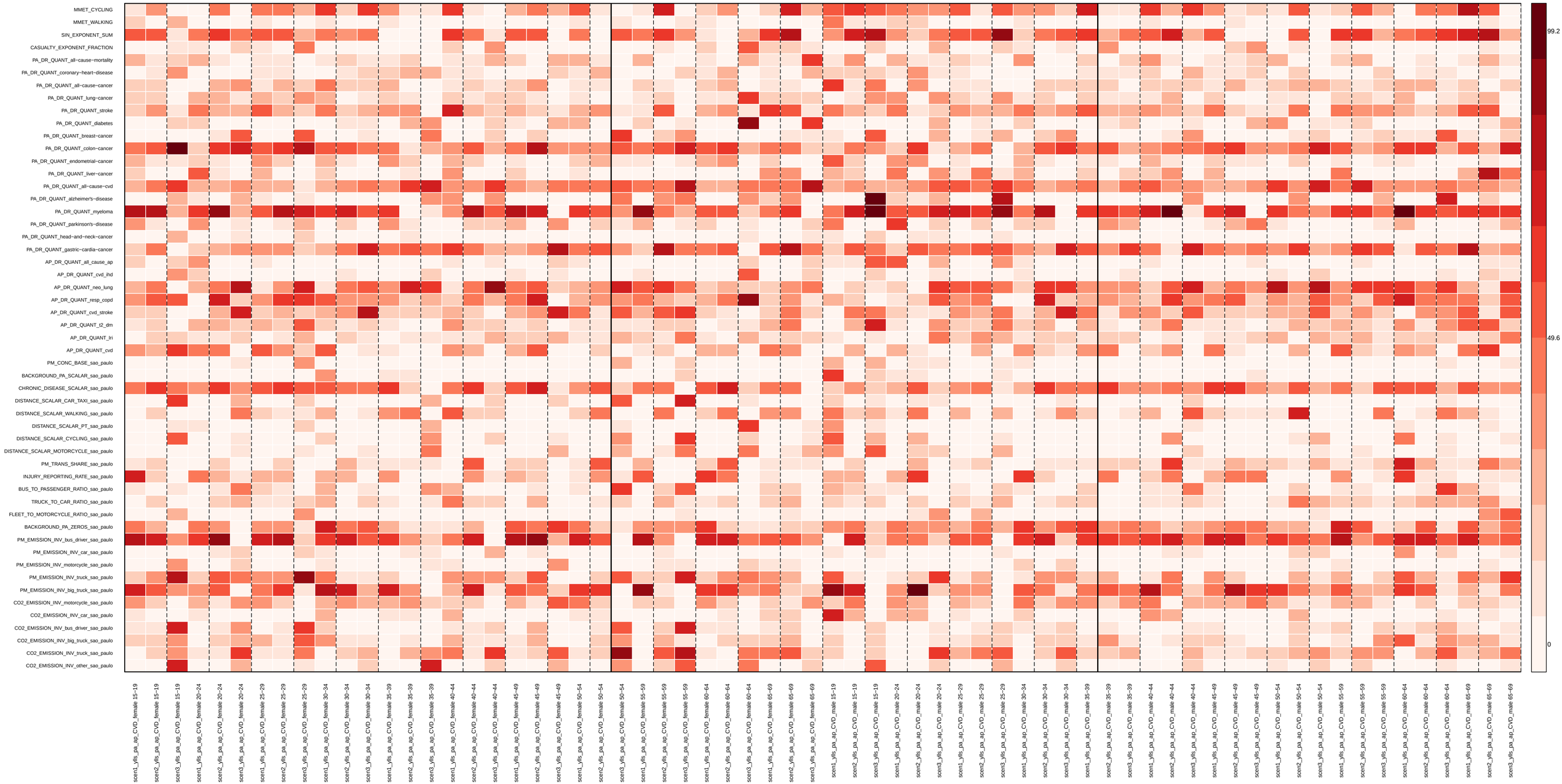
sao_paulo − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



sao_paulo − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

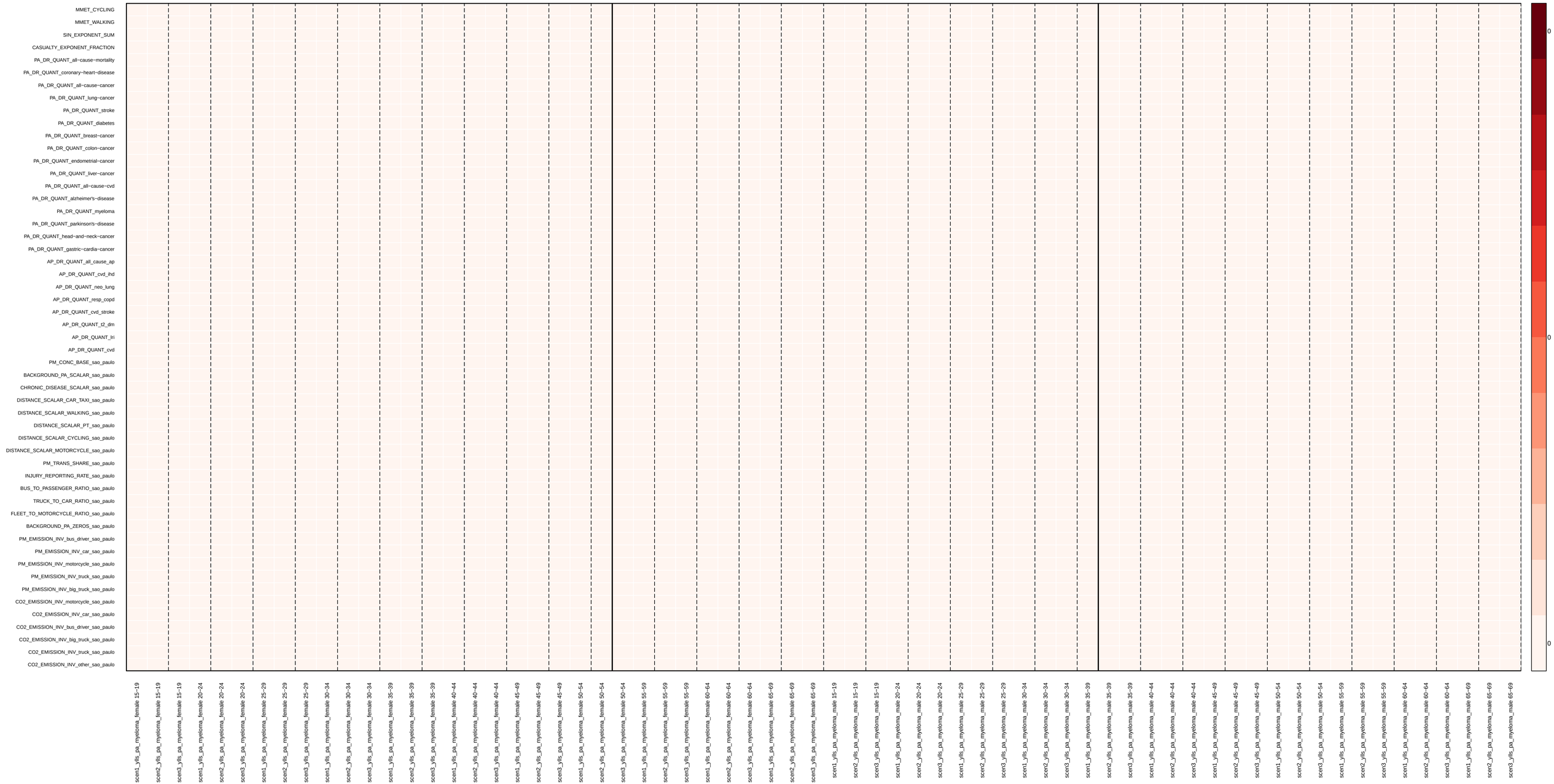


sao_paulo − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

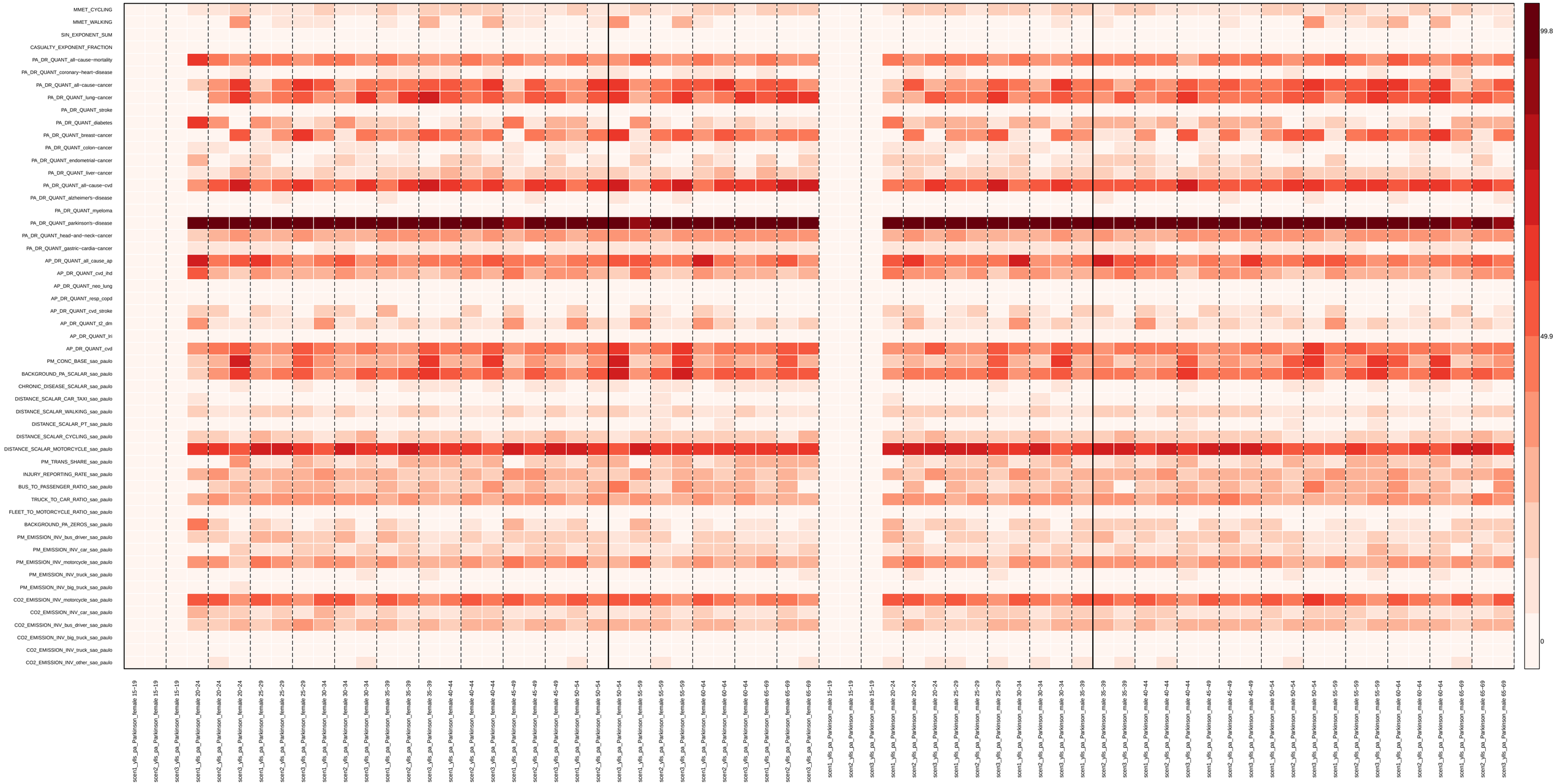


sao_paulo − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

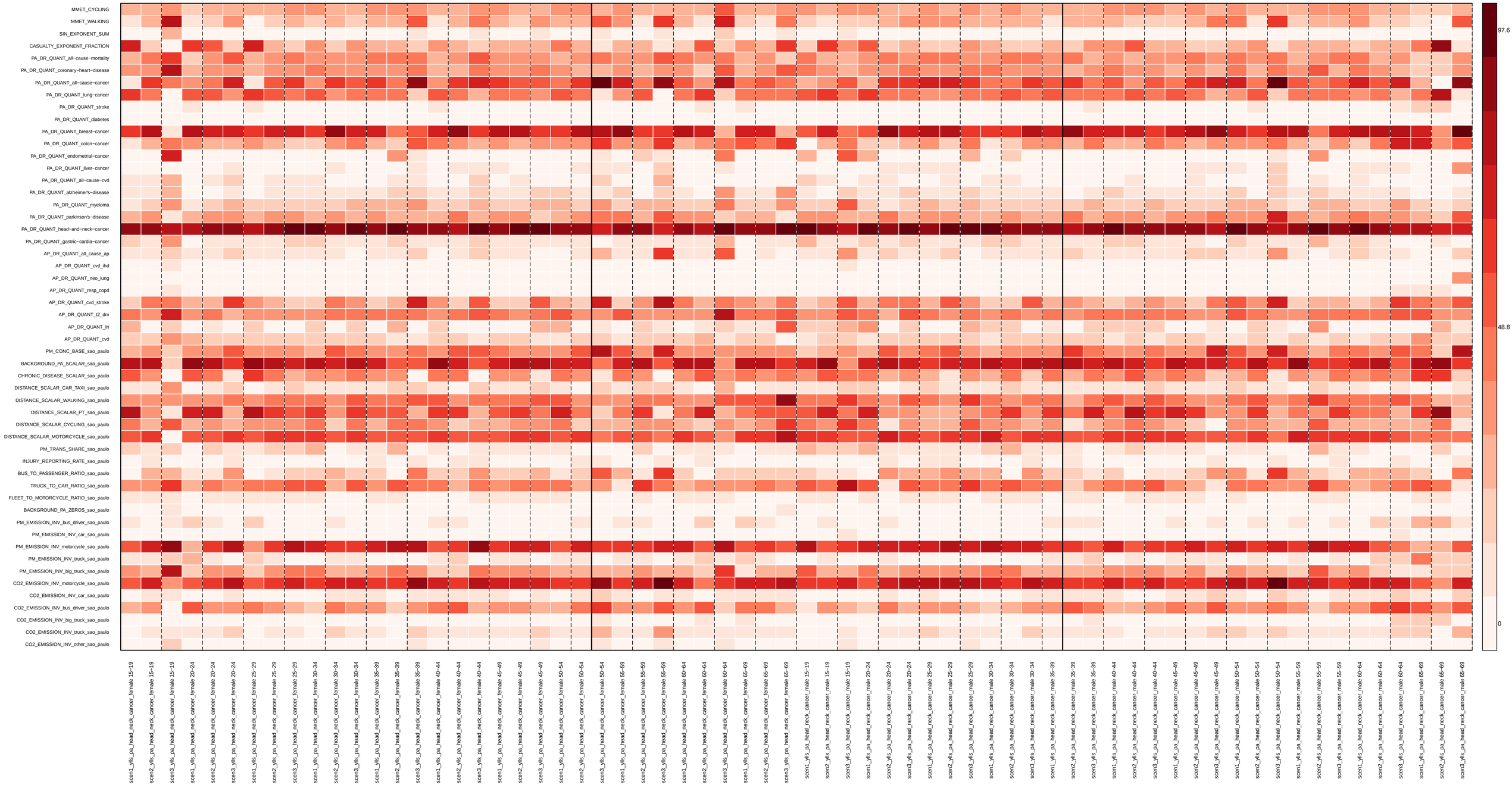
sao_paulo − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



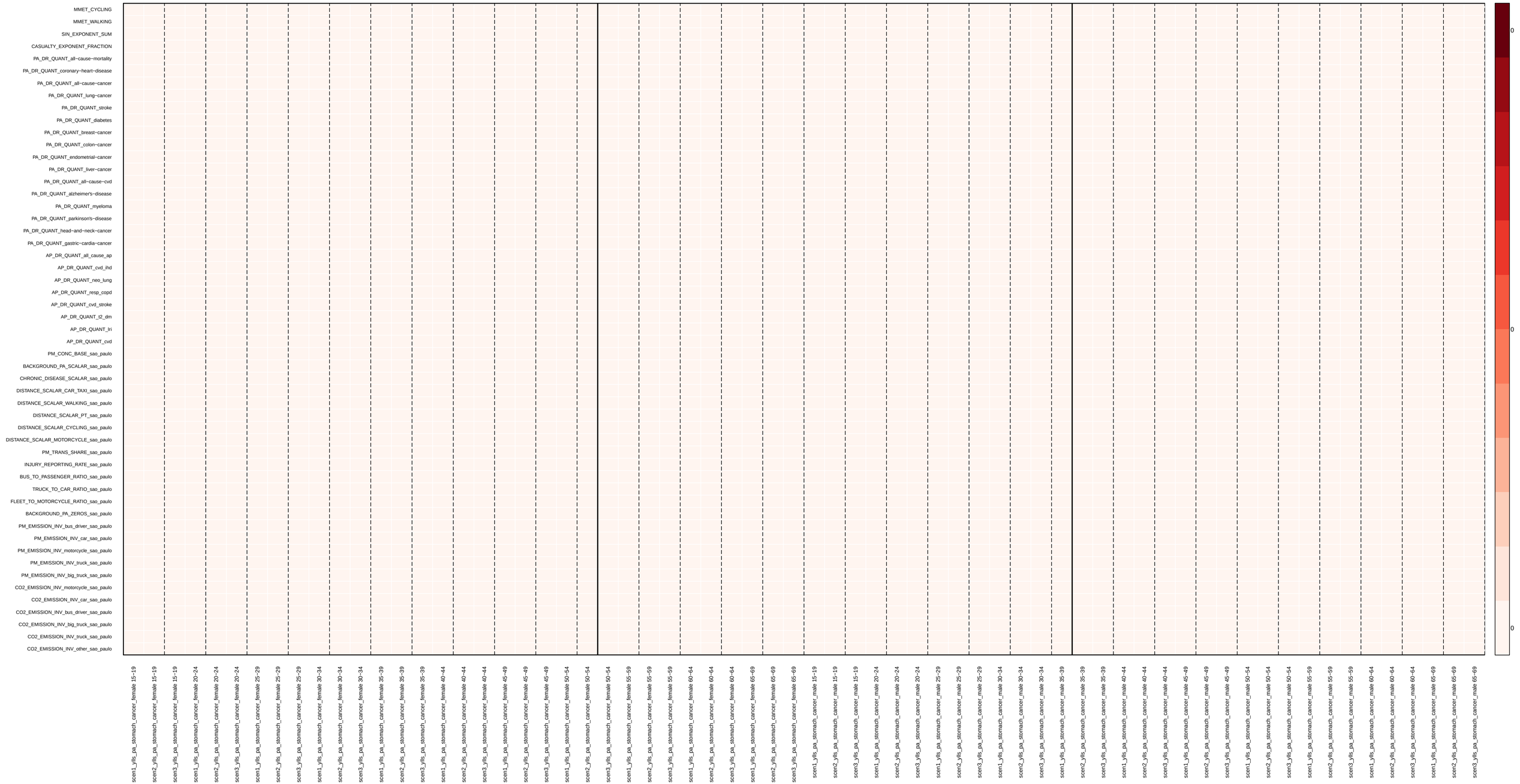
sao_paulo − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



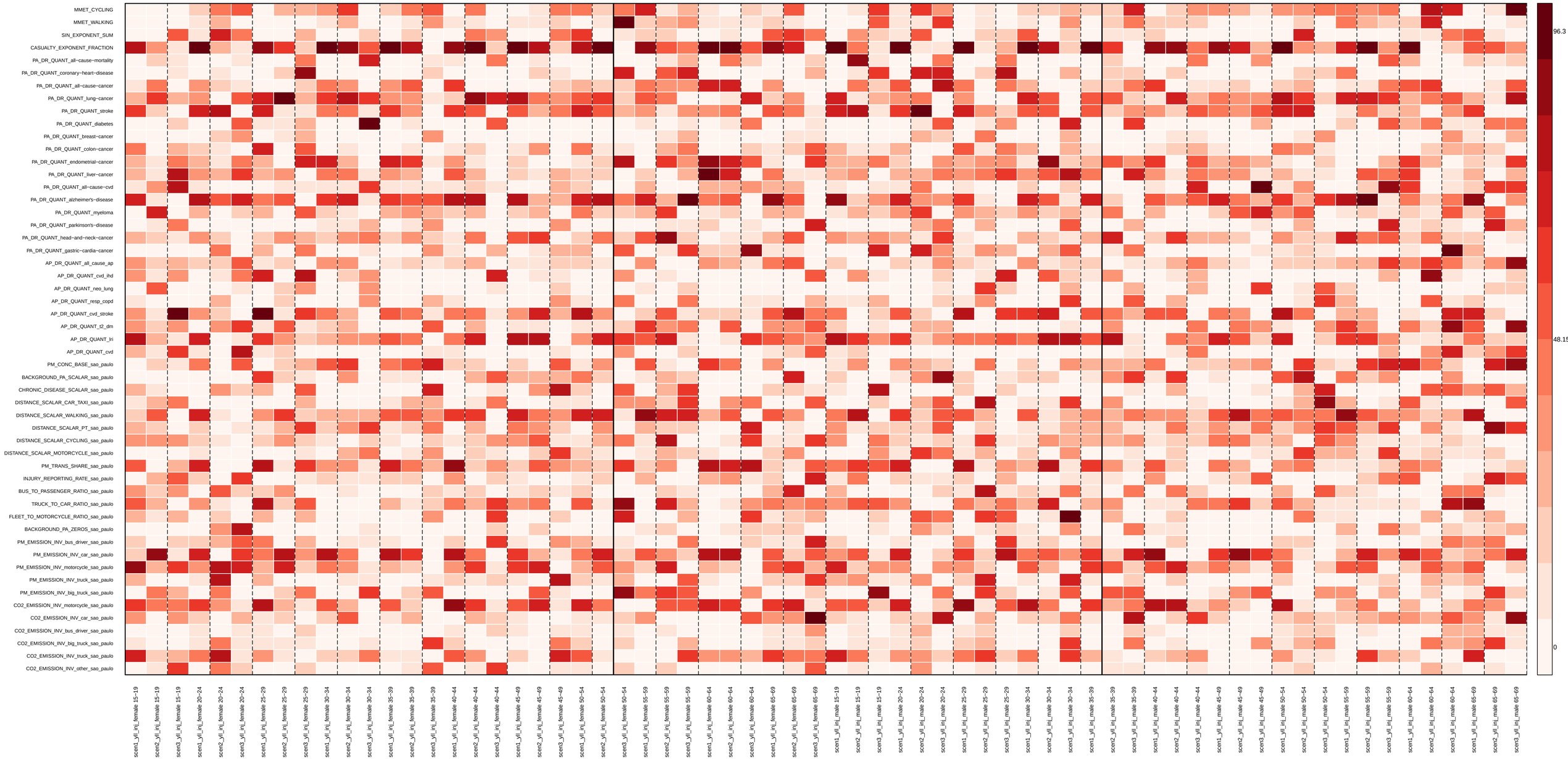
sao_paulo − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



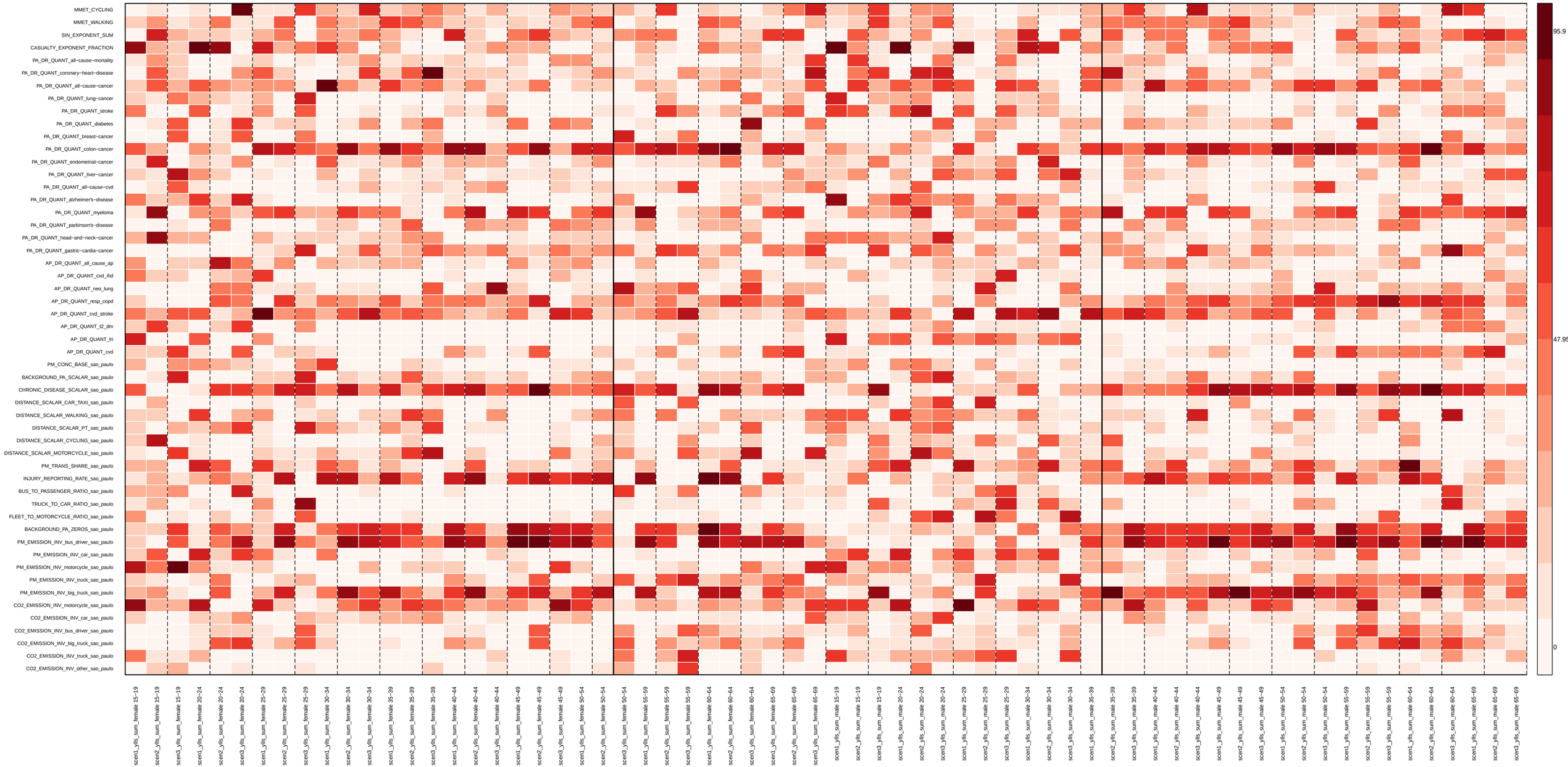
sao_paulo − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



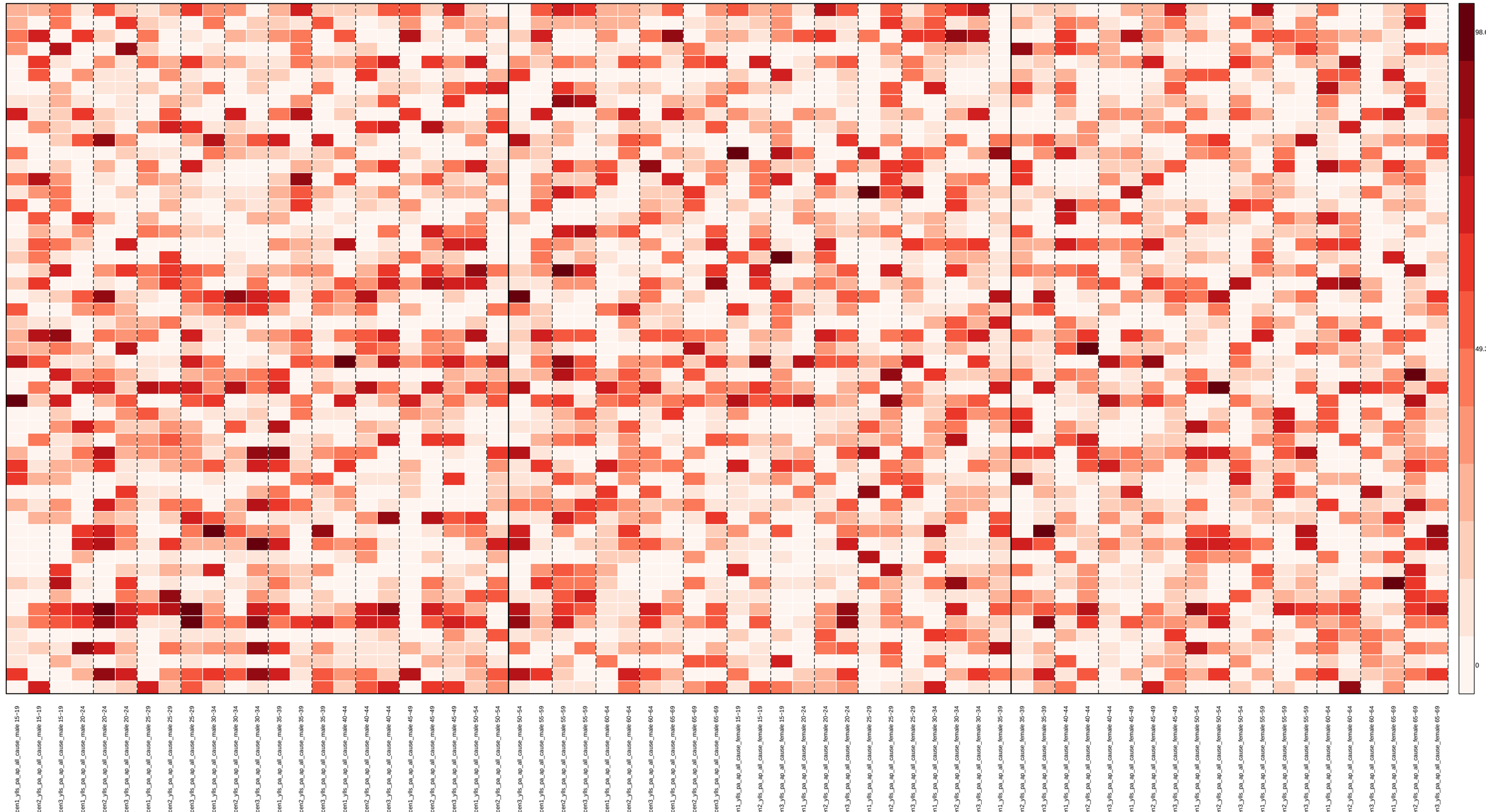
sao_paulo − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



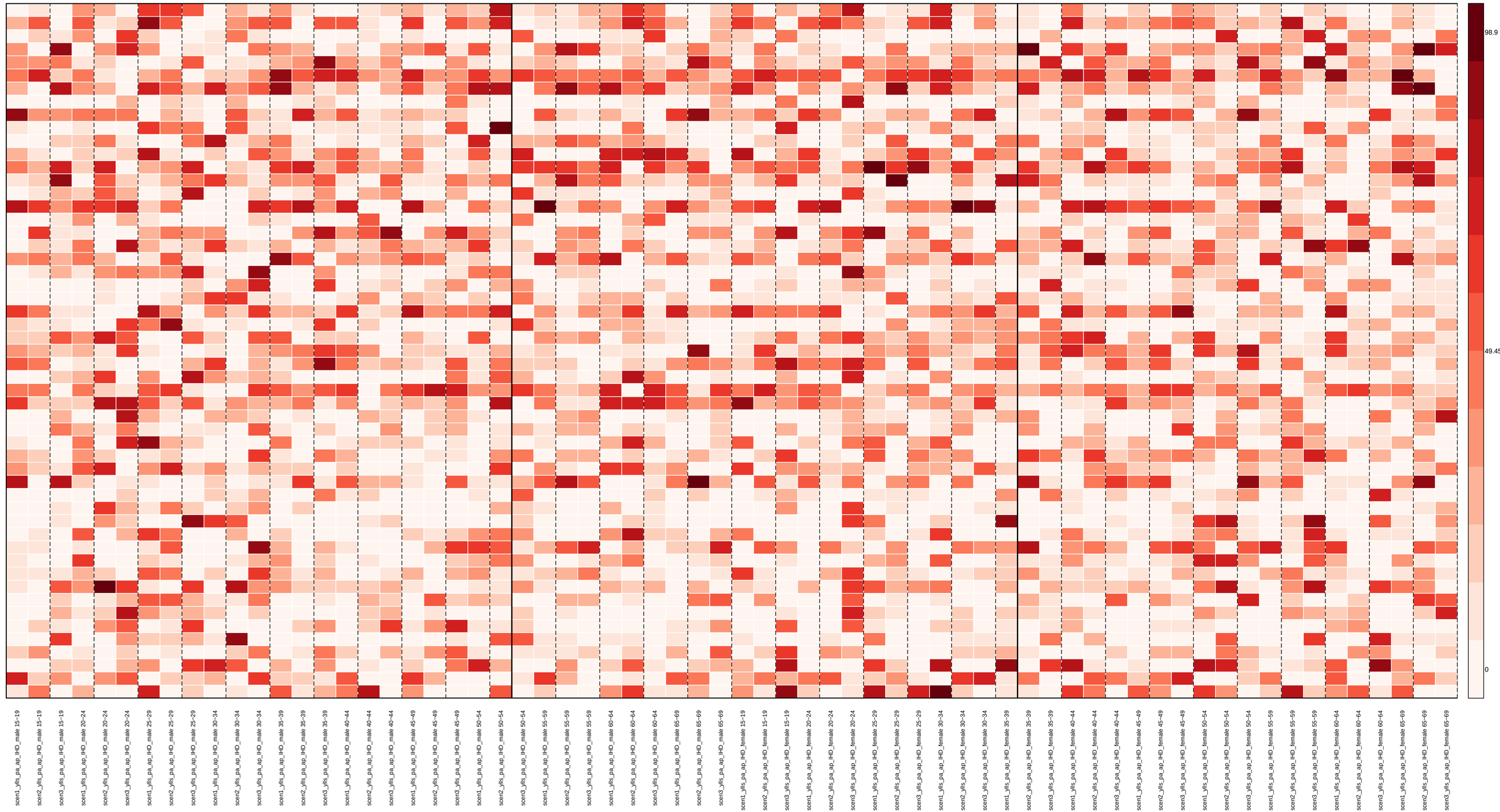
sao_paulo − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



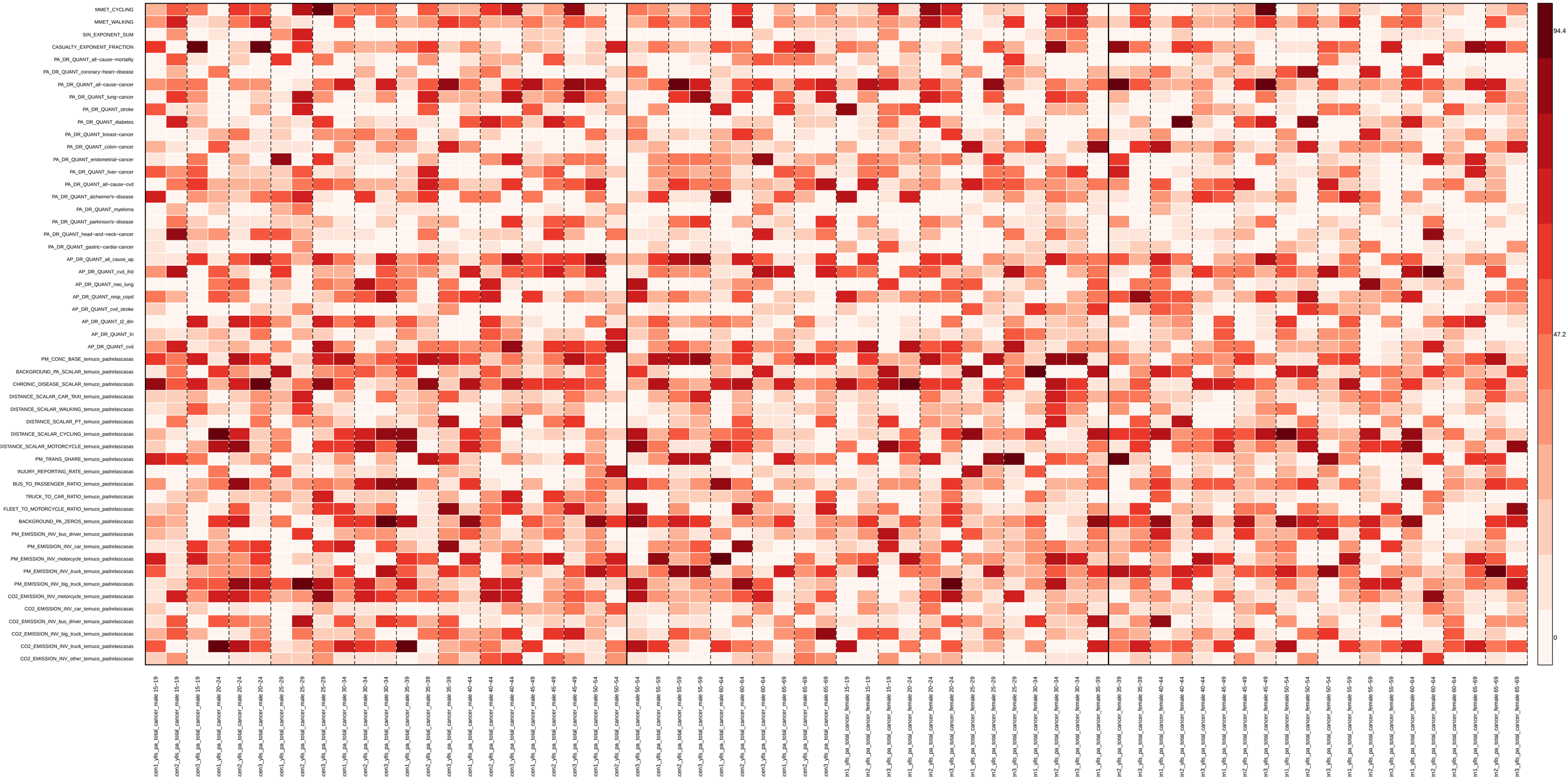
temuco_padrelascasas − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



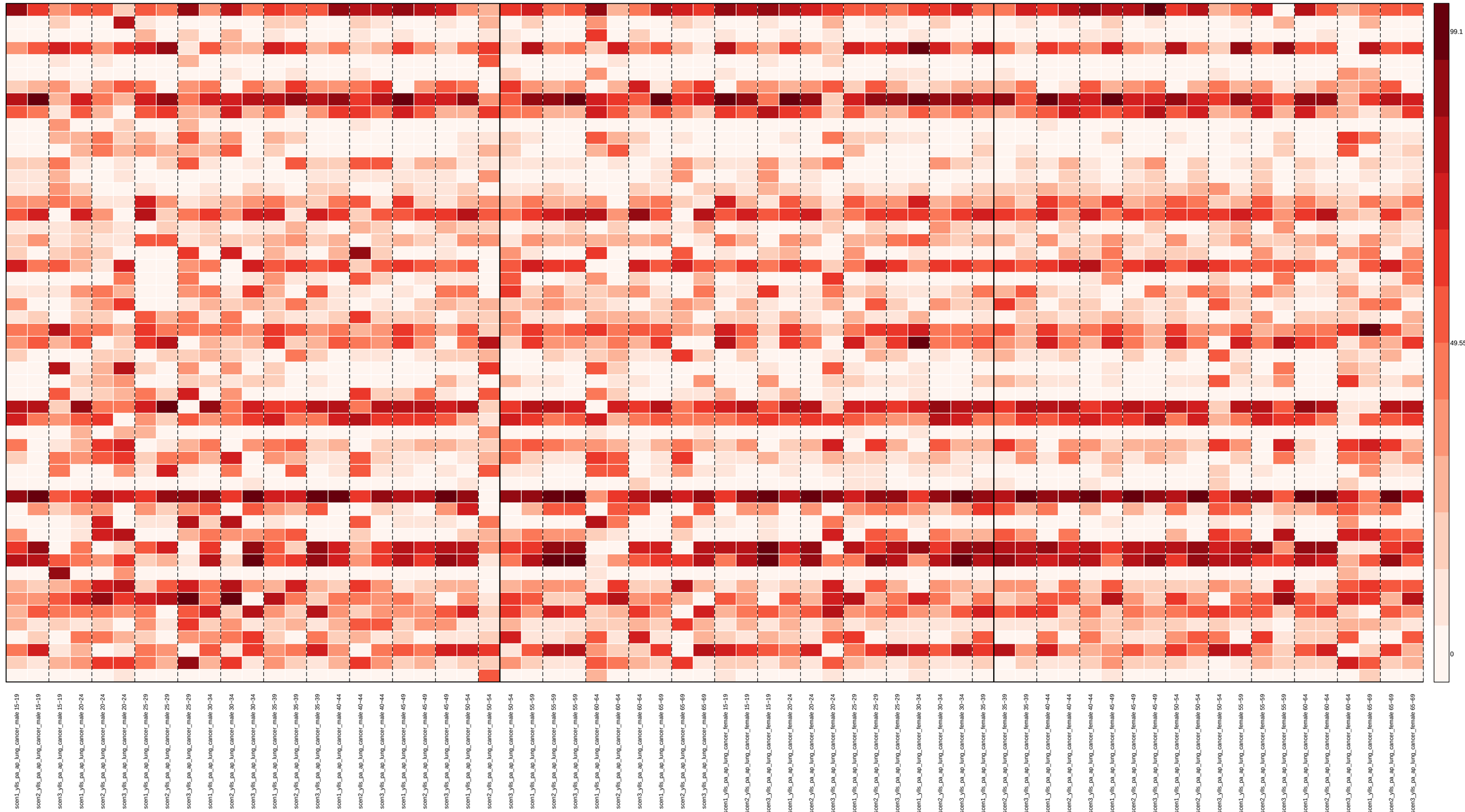
temuco_padrelascasas − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



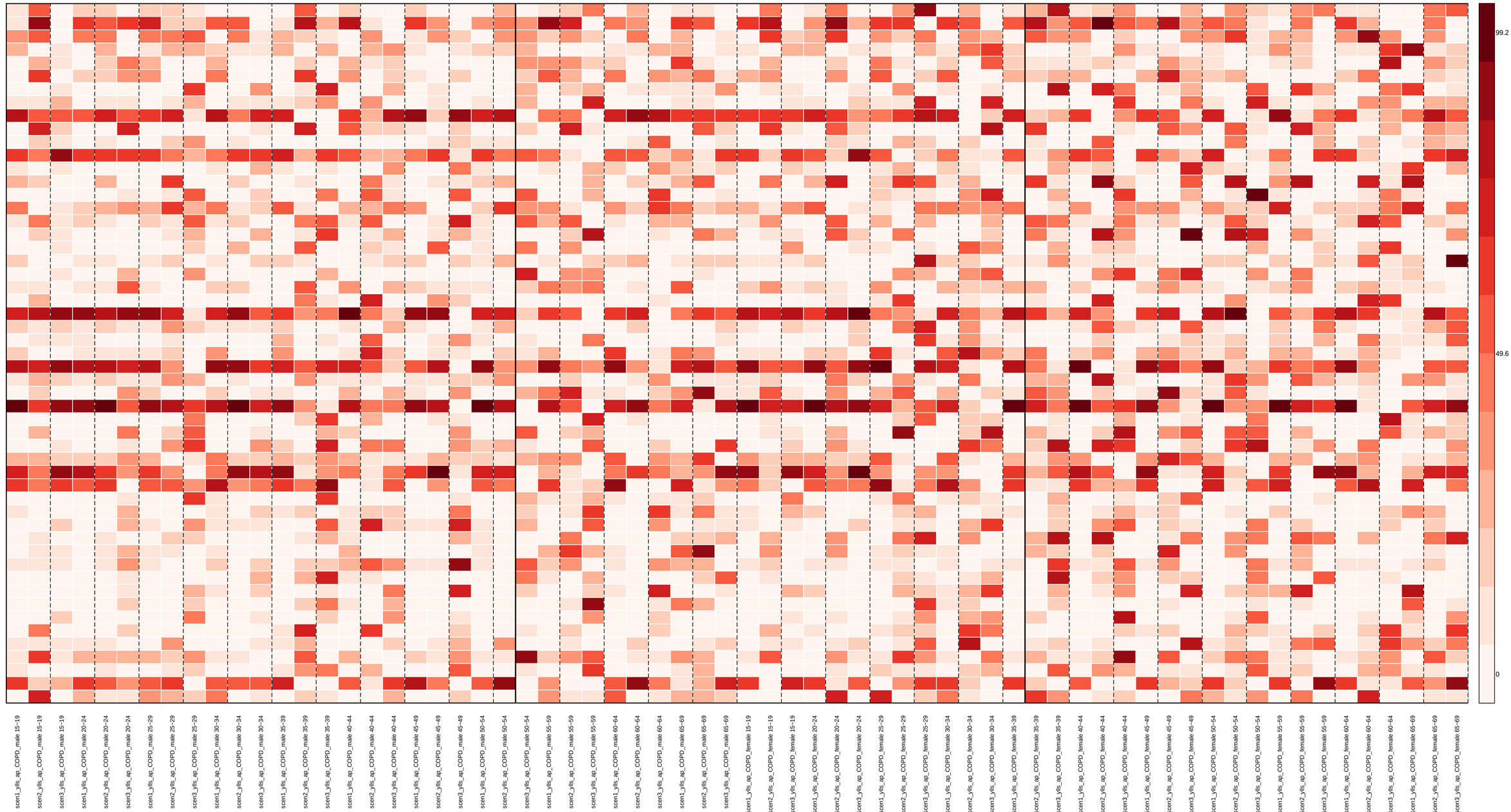
temuco_padrelascasas − pa_total_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



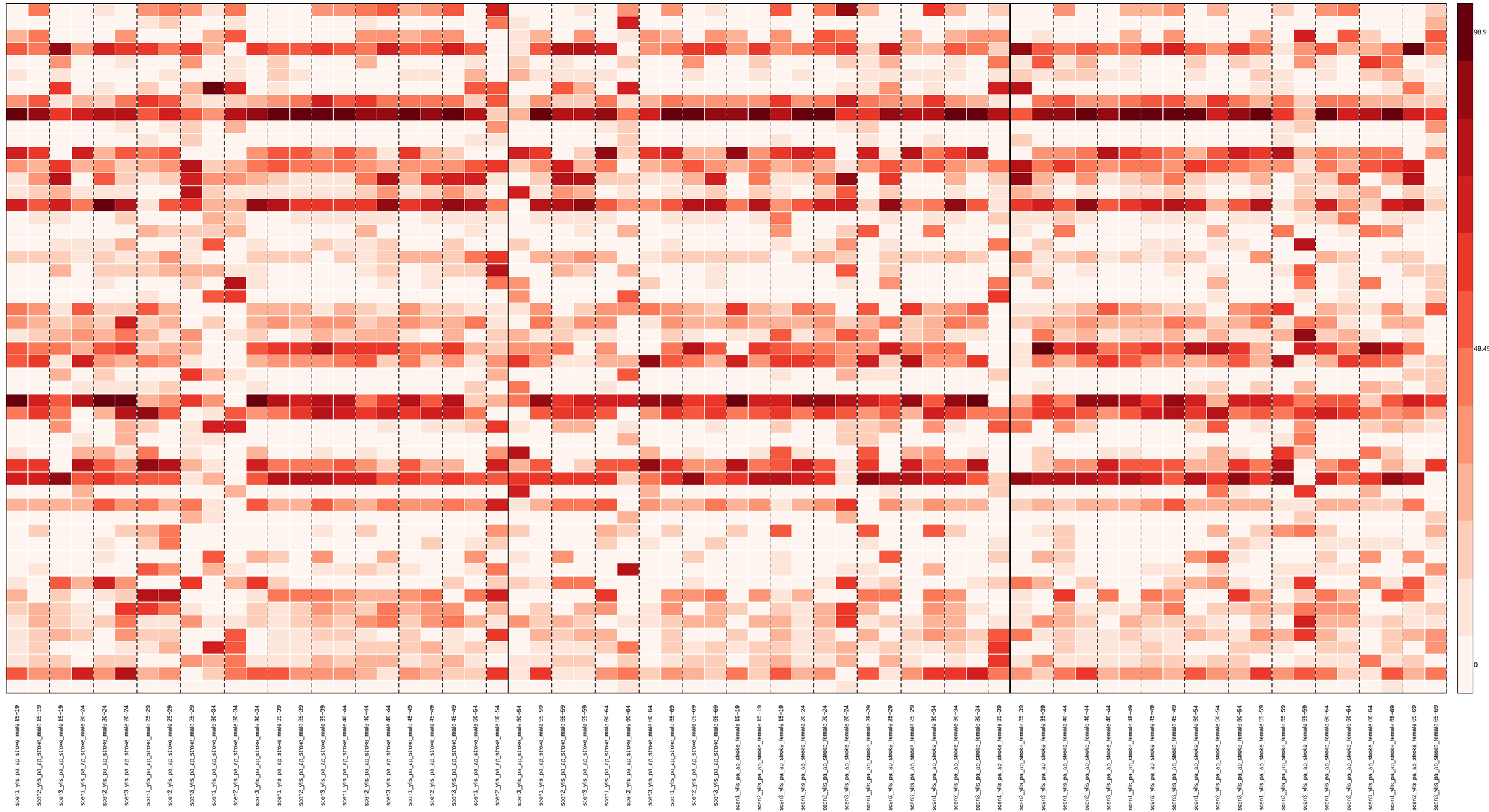
temuco_padrelascasas − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



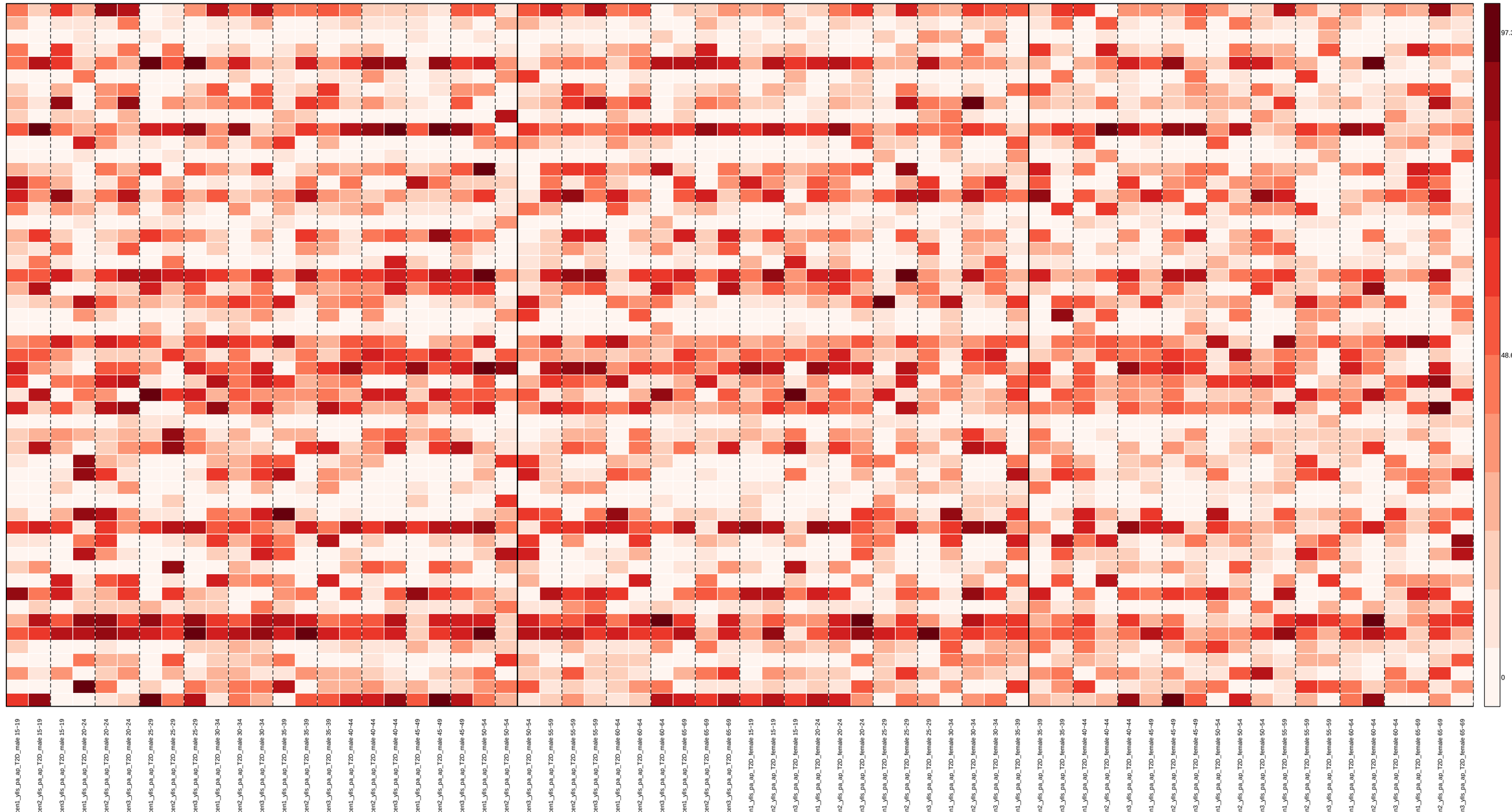
temuco_padrelascasas − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



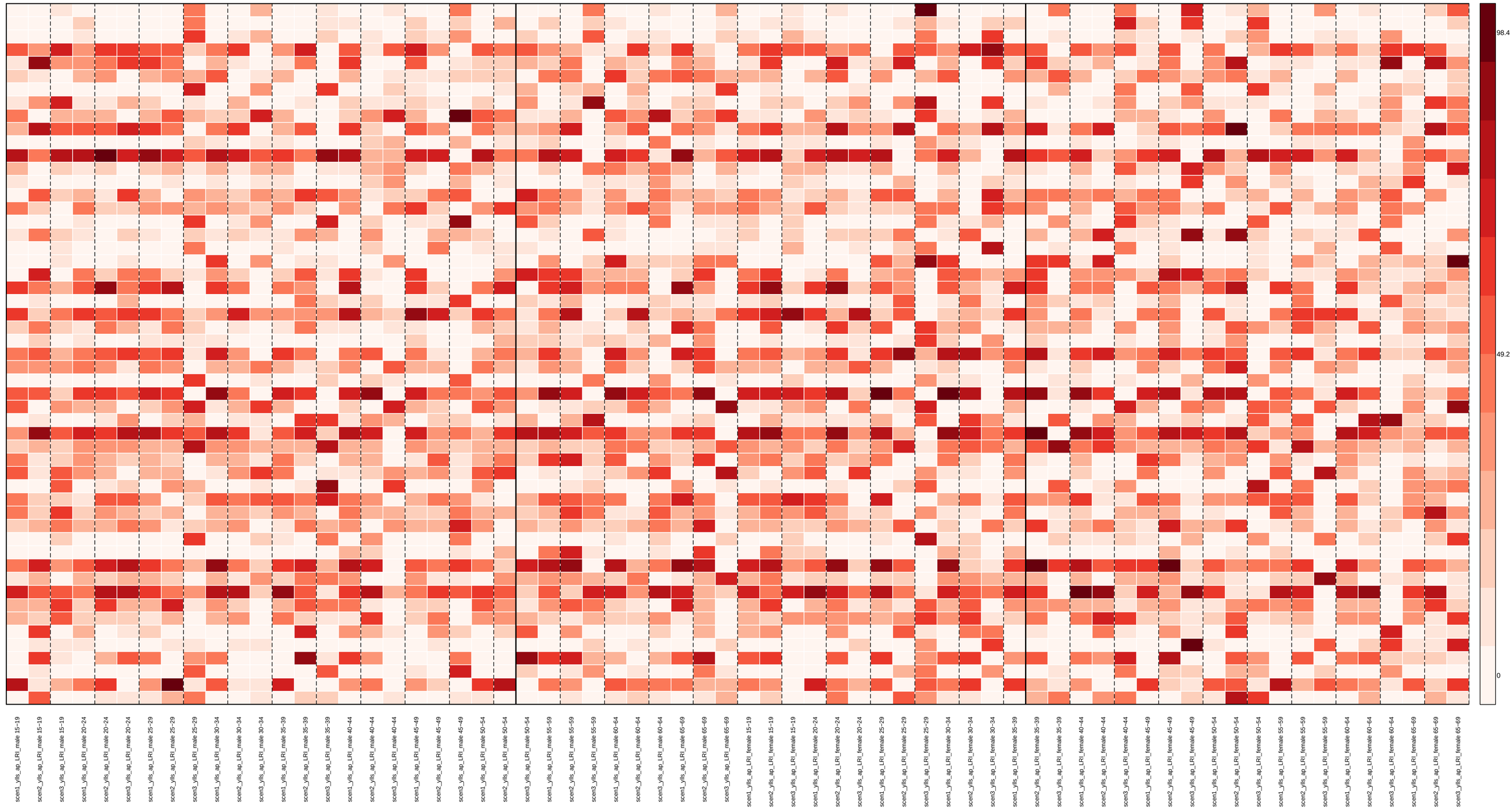
temuco_padrelascasas − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



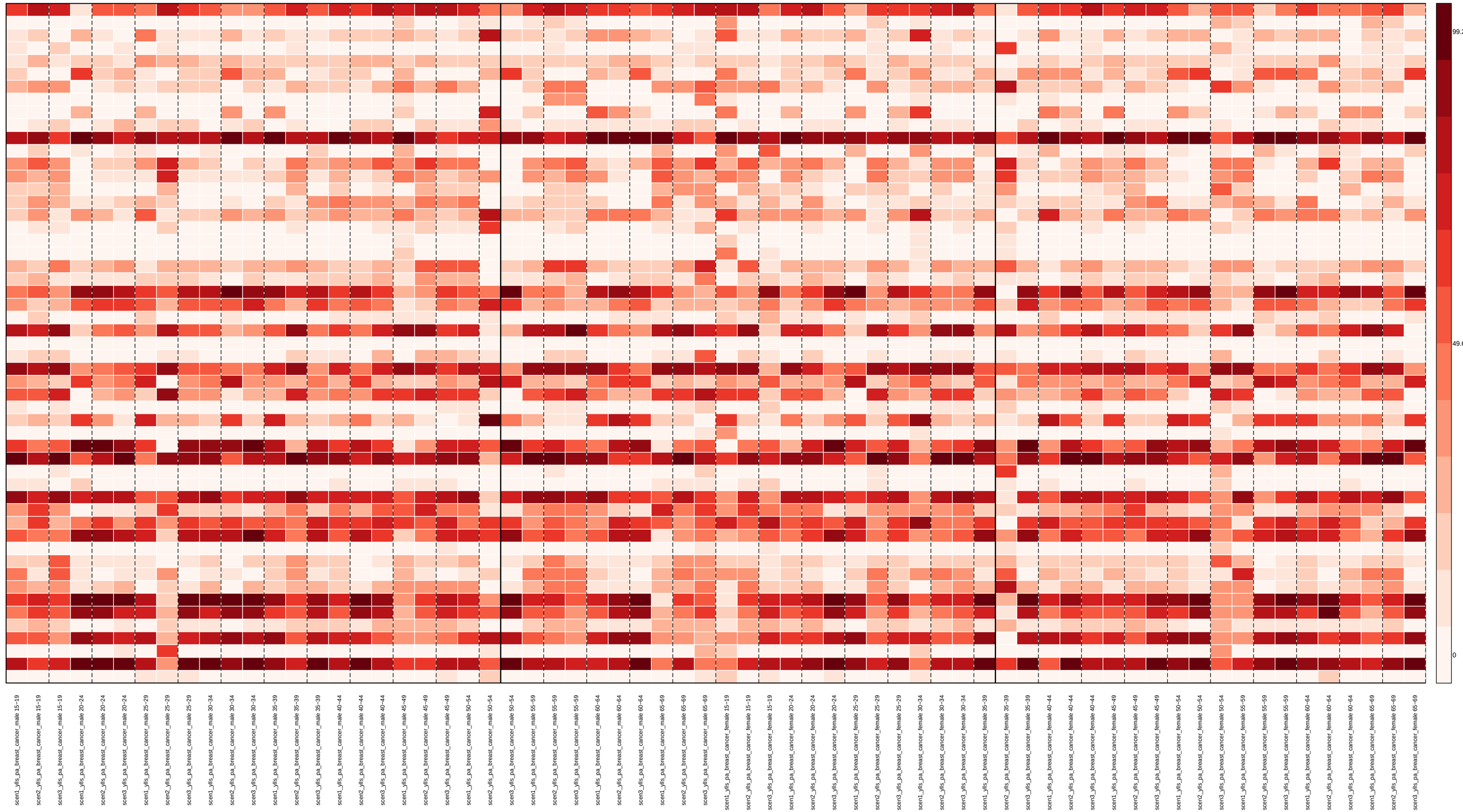
temuco_padrelascasas − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



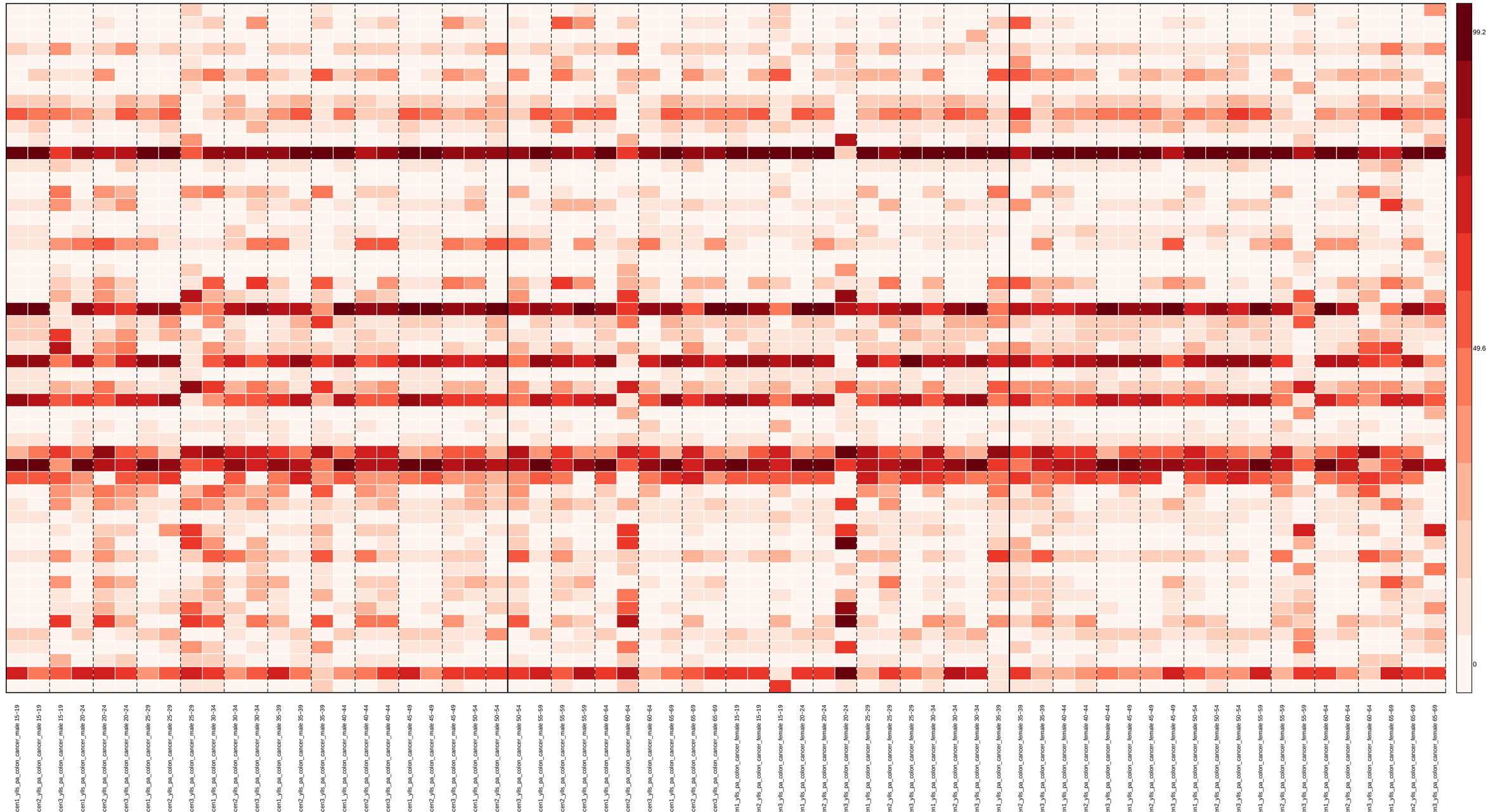
temuco_padrelascasas − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



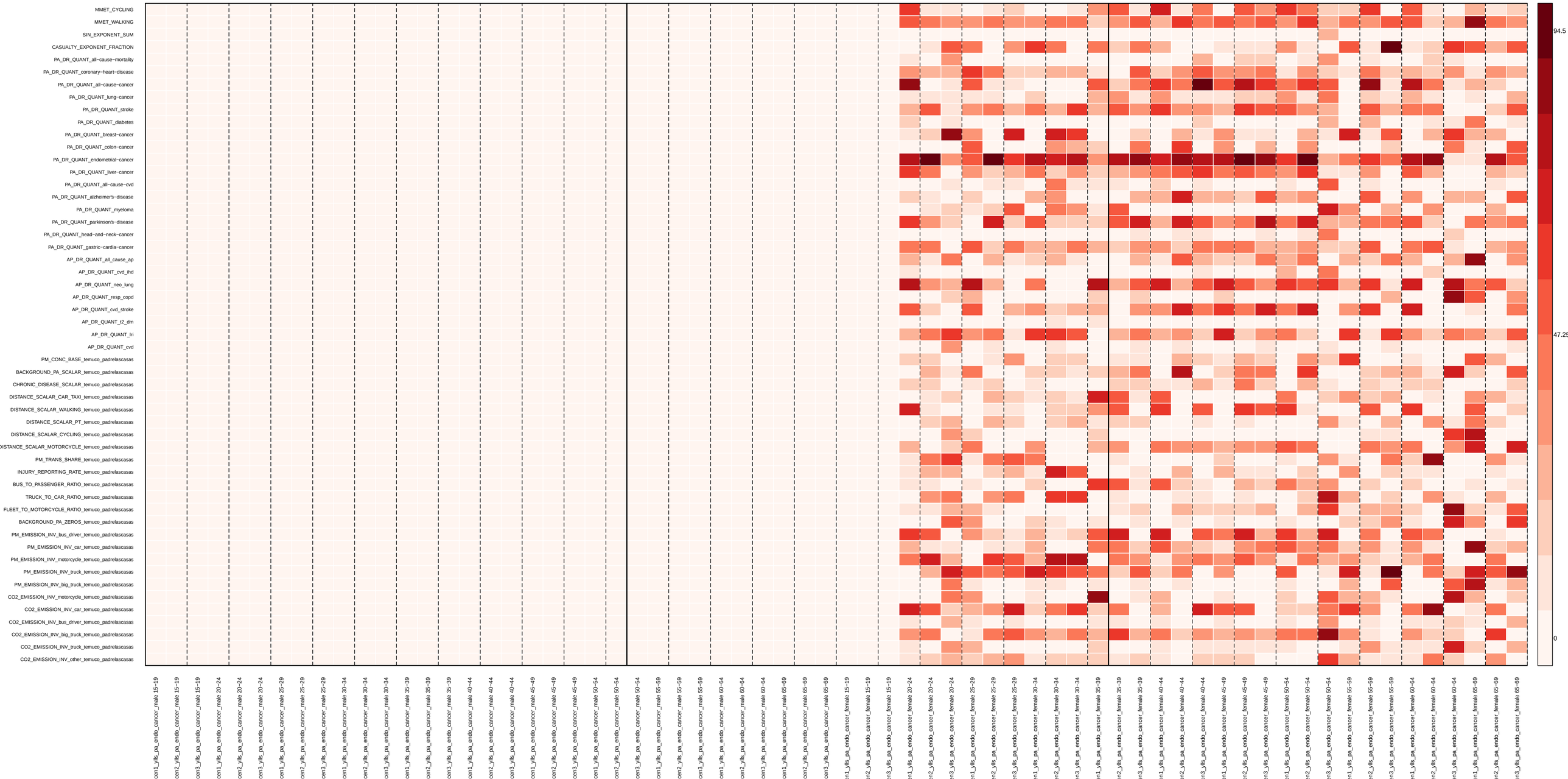
temuco_padrelascasas − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



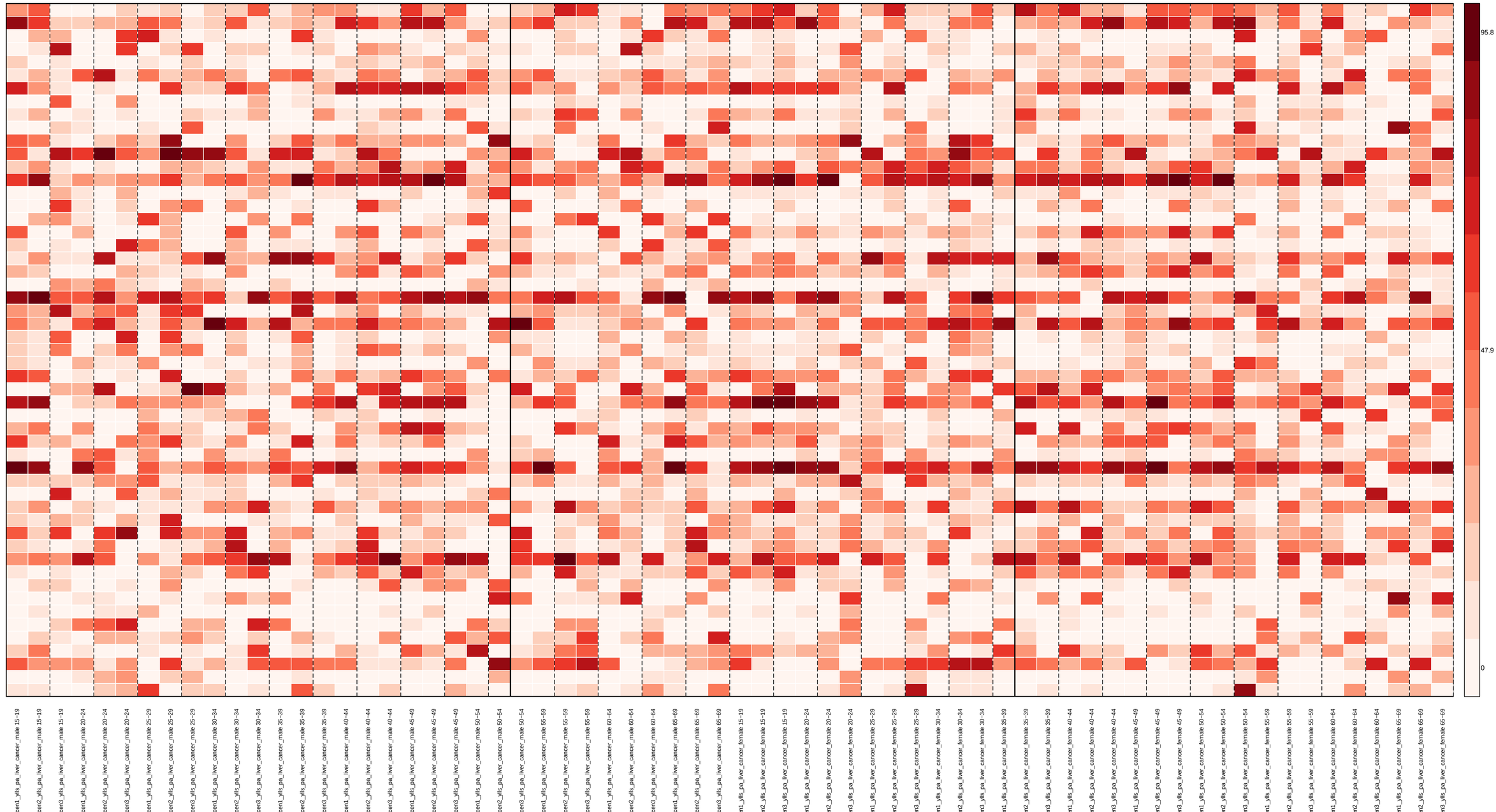
temuco_padrelascasas − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



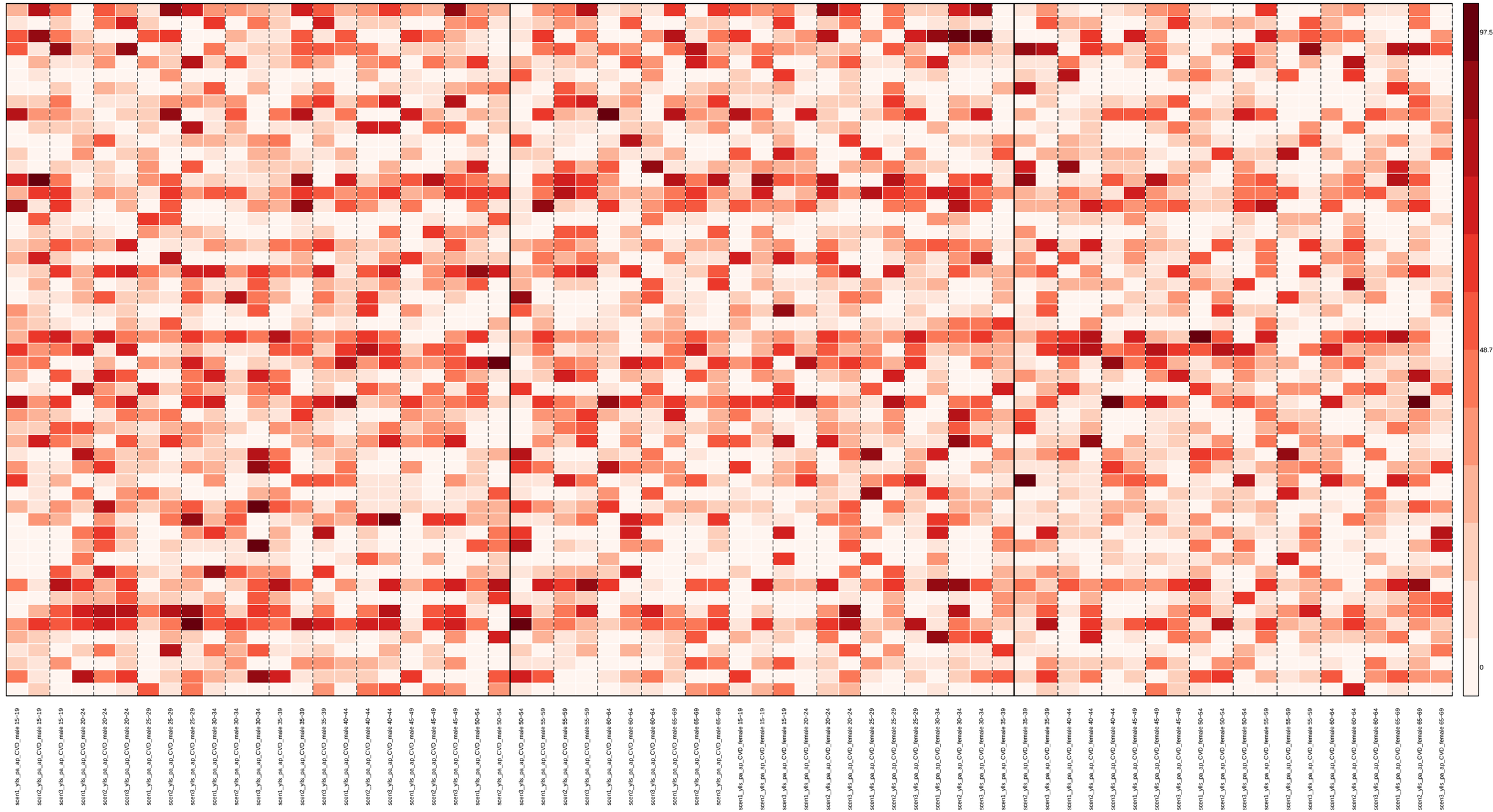
temuco_padrelascasas − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



temuco_padrelascasas − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

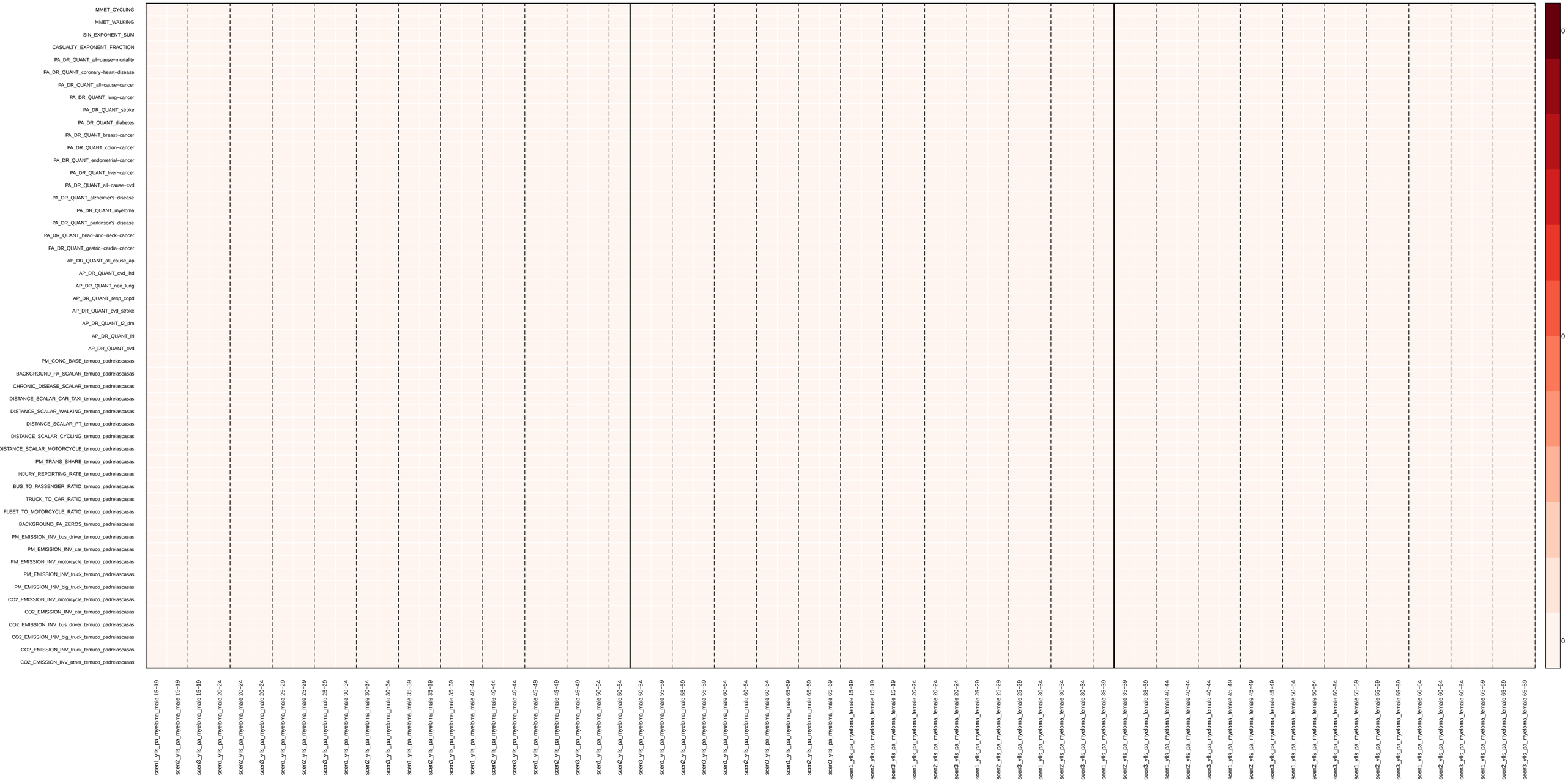


temuco_padrelascasas − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

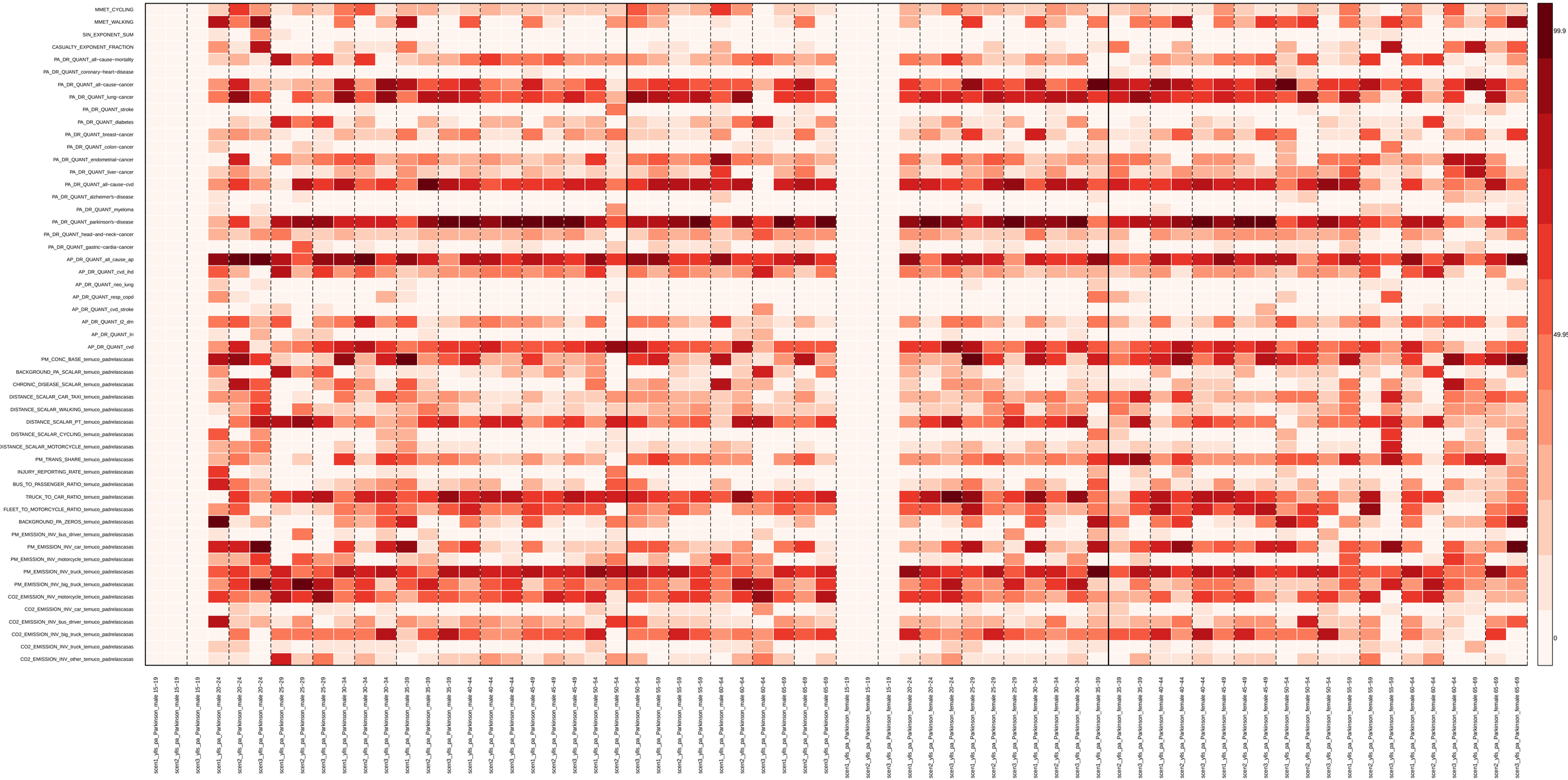


temuco_padrelascasas − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?

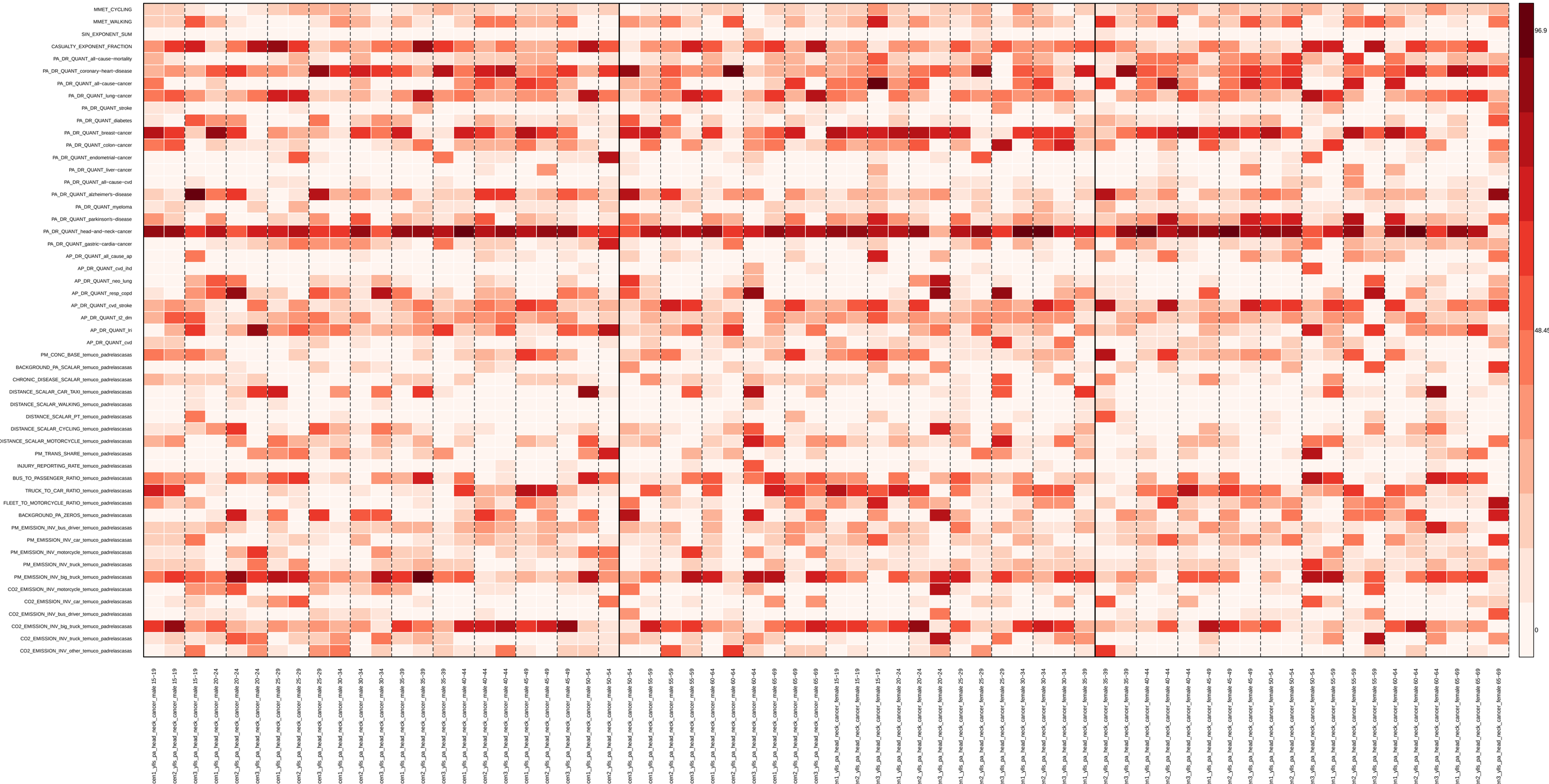
temuco_padrelascasas − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



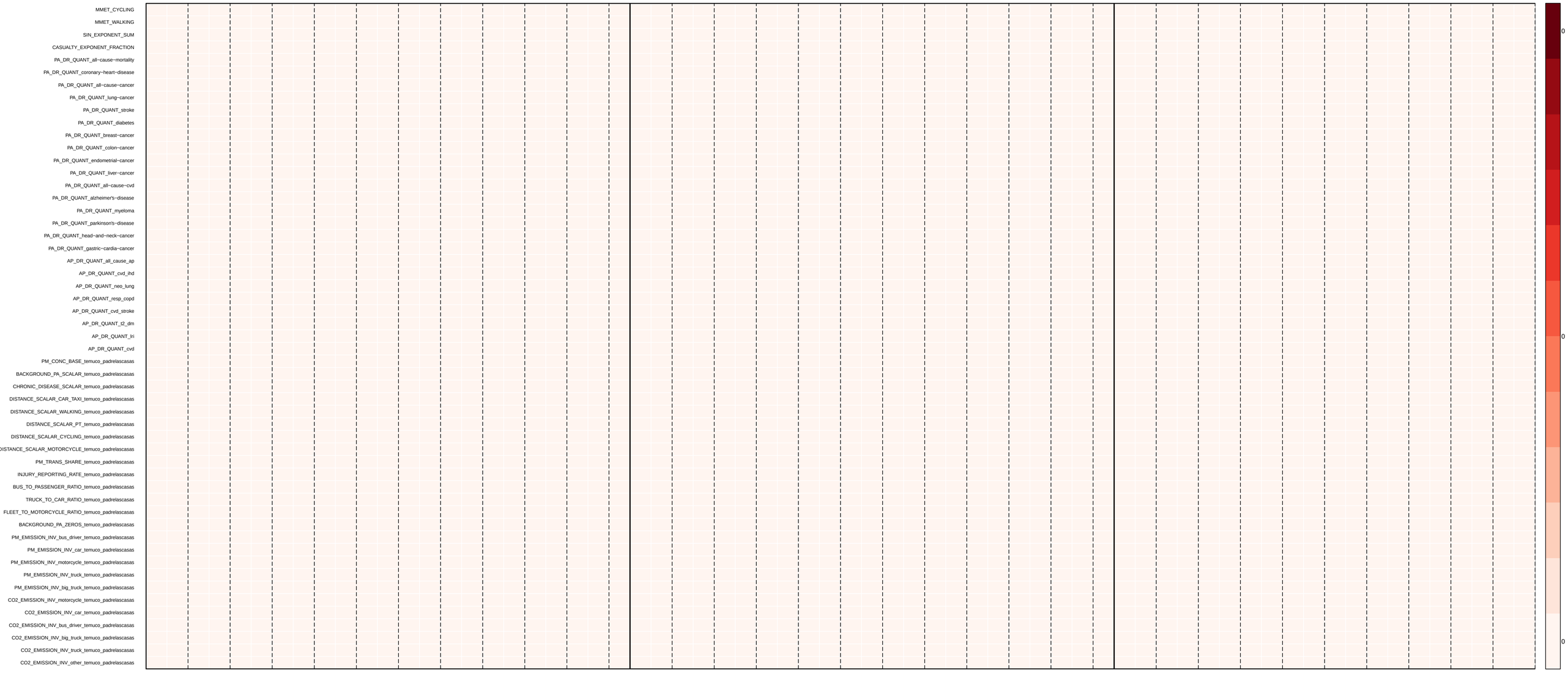
temuco_padrelascasas − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



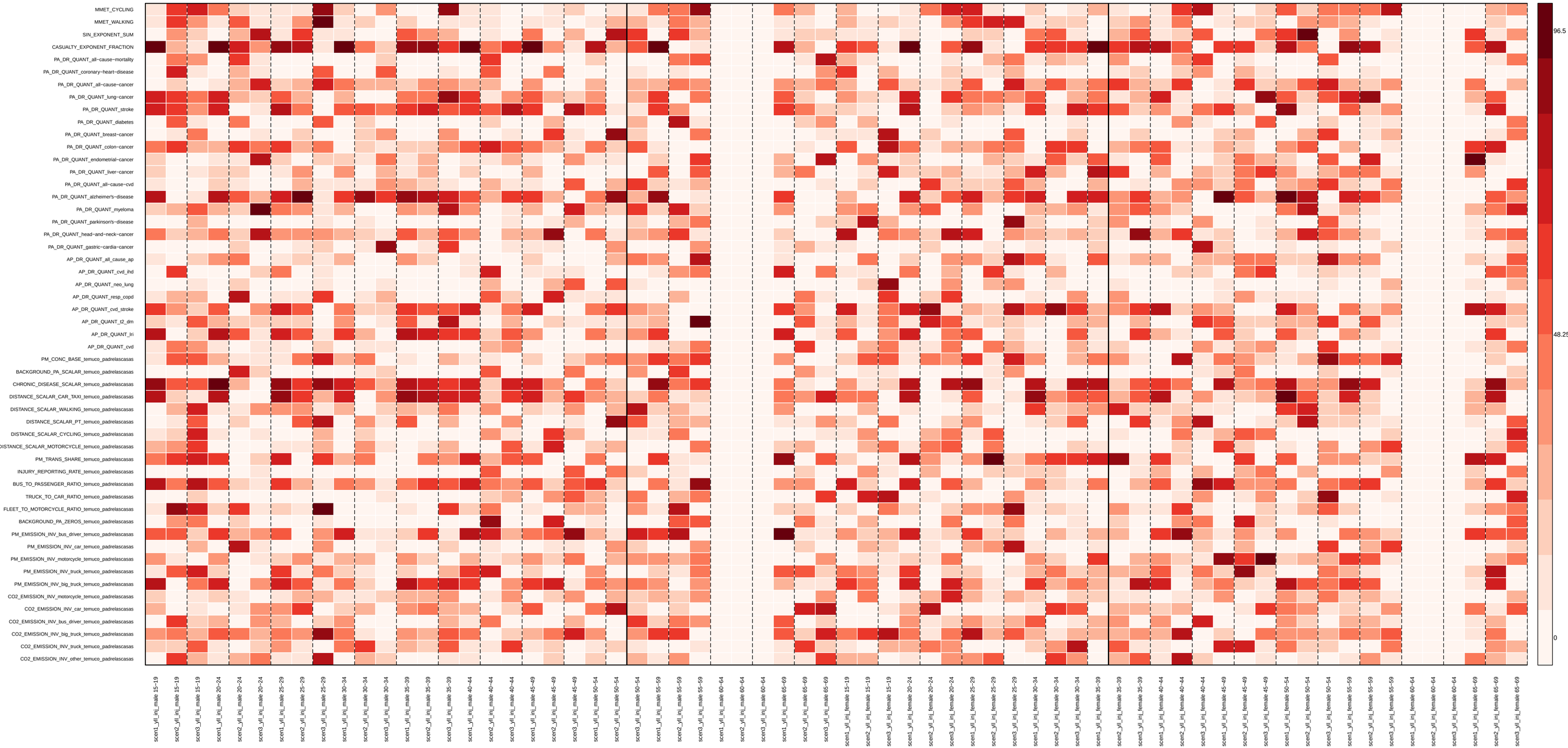
temuco_padrelascasas − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



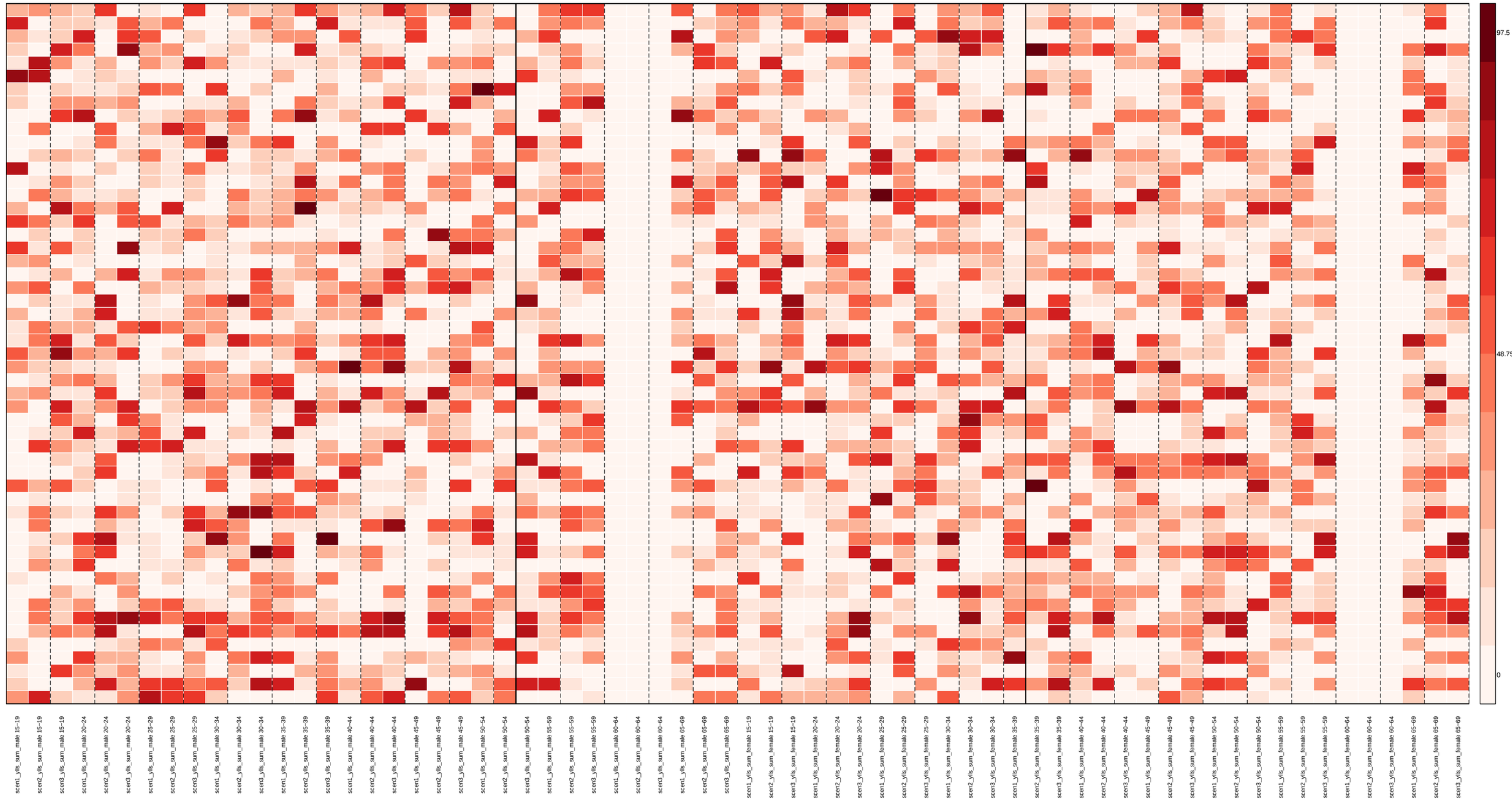
temuco_padrelascasas − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



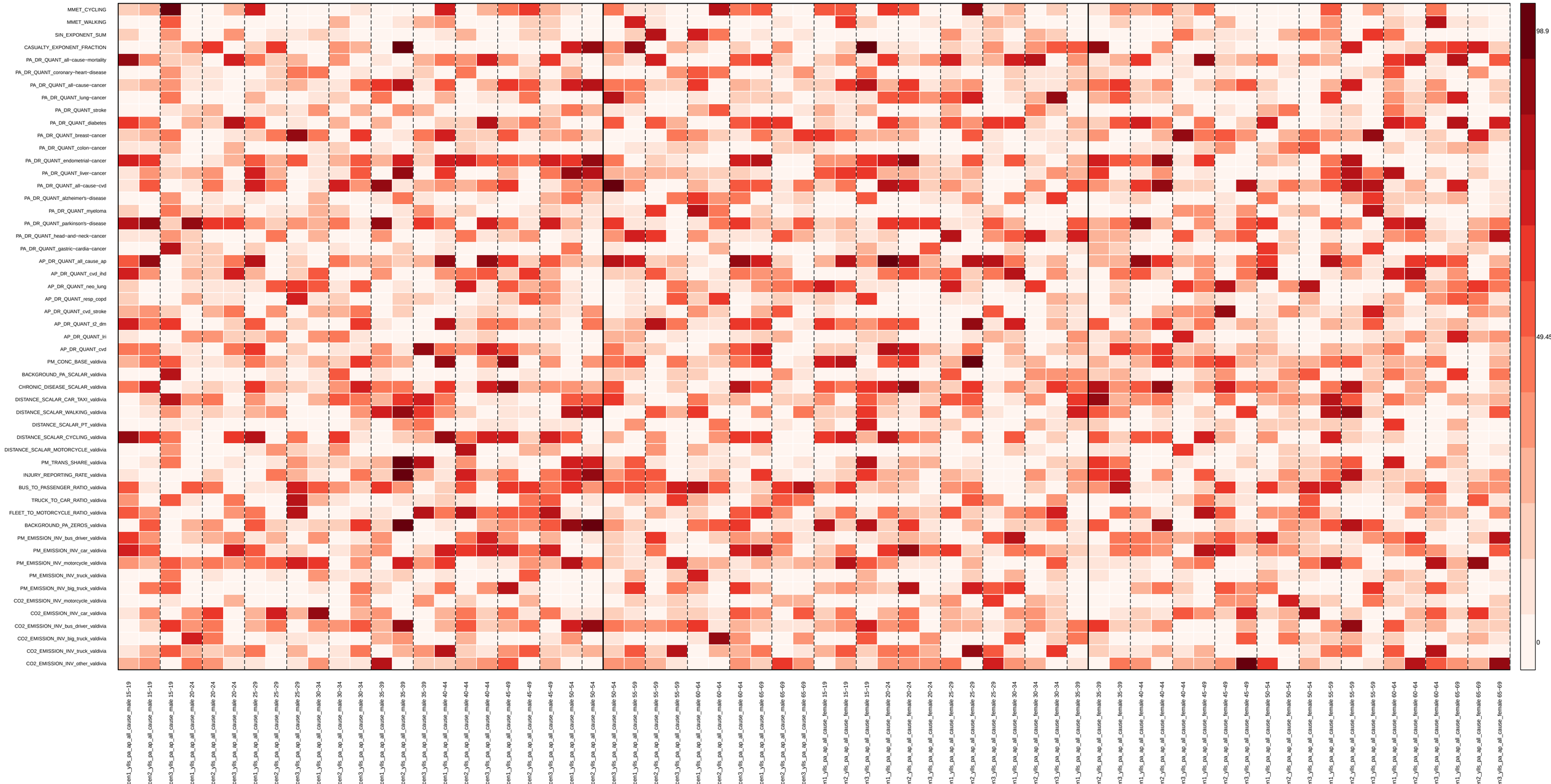
temuco_padrelascasas − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



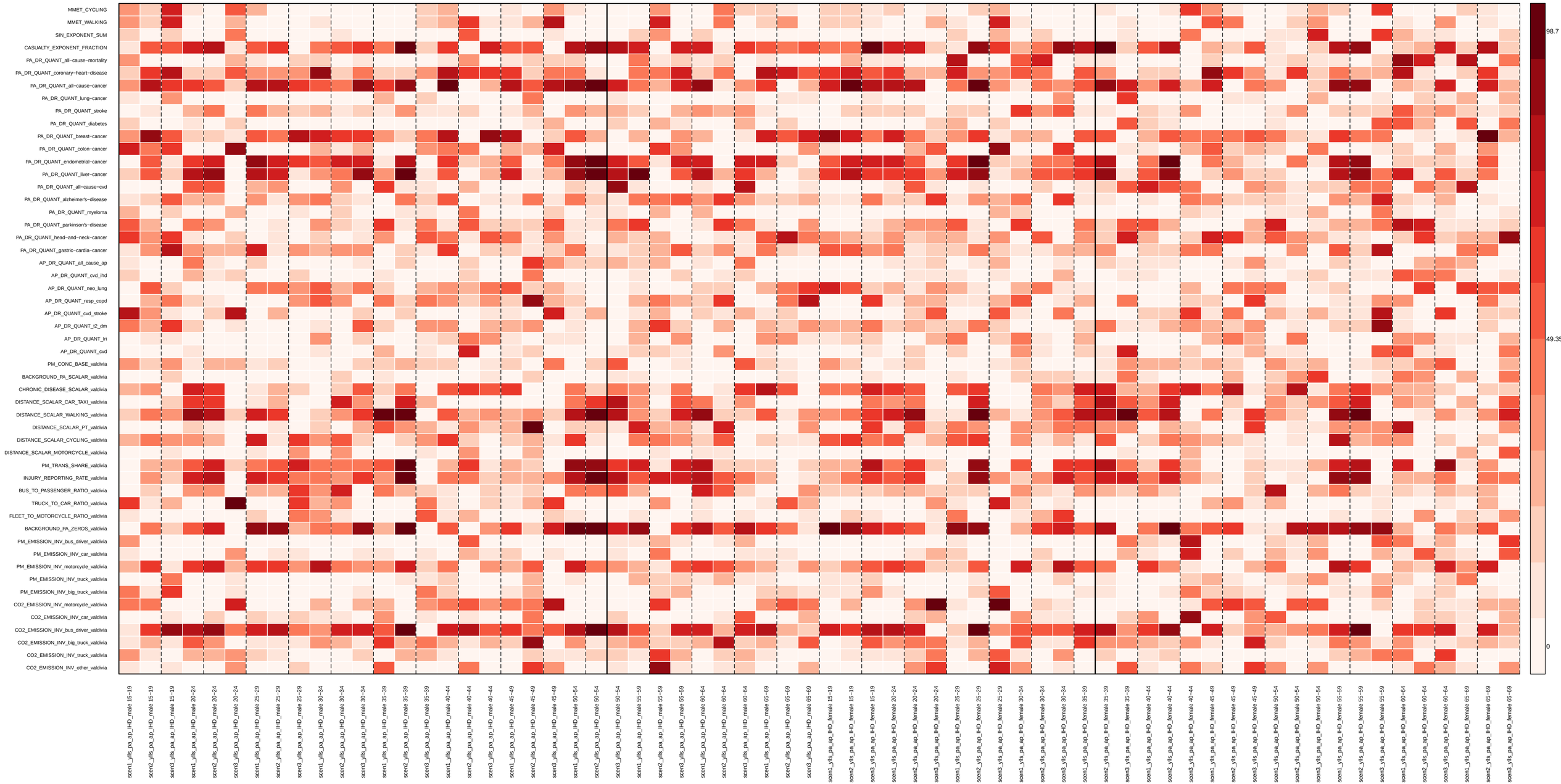
temuco_padrelascasas − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



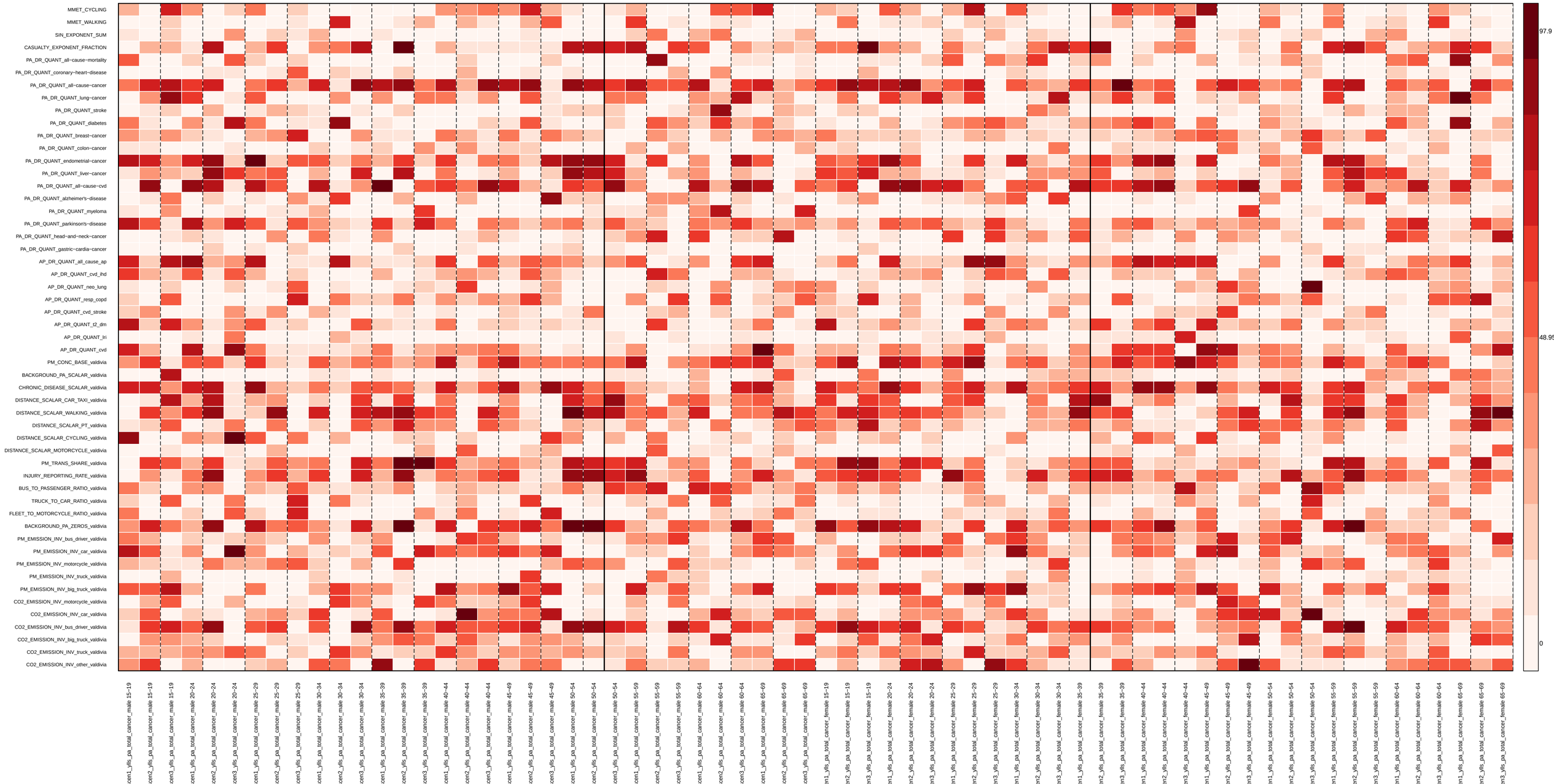
valdivia − pa_ap_all_cause , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



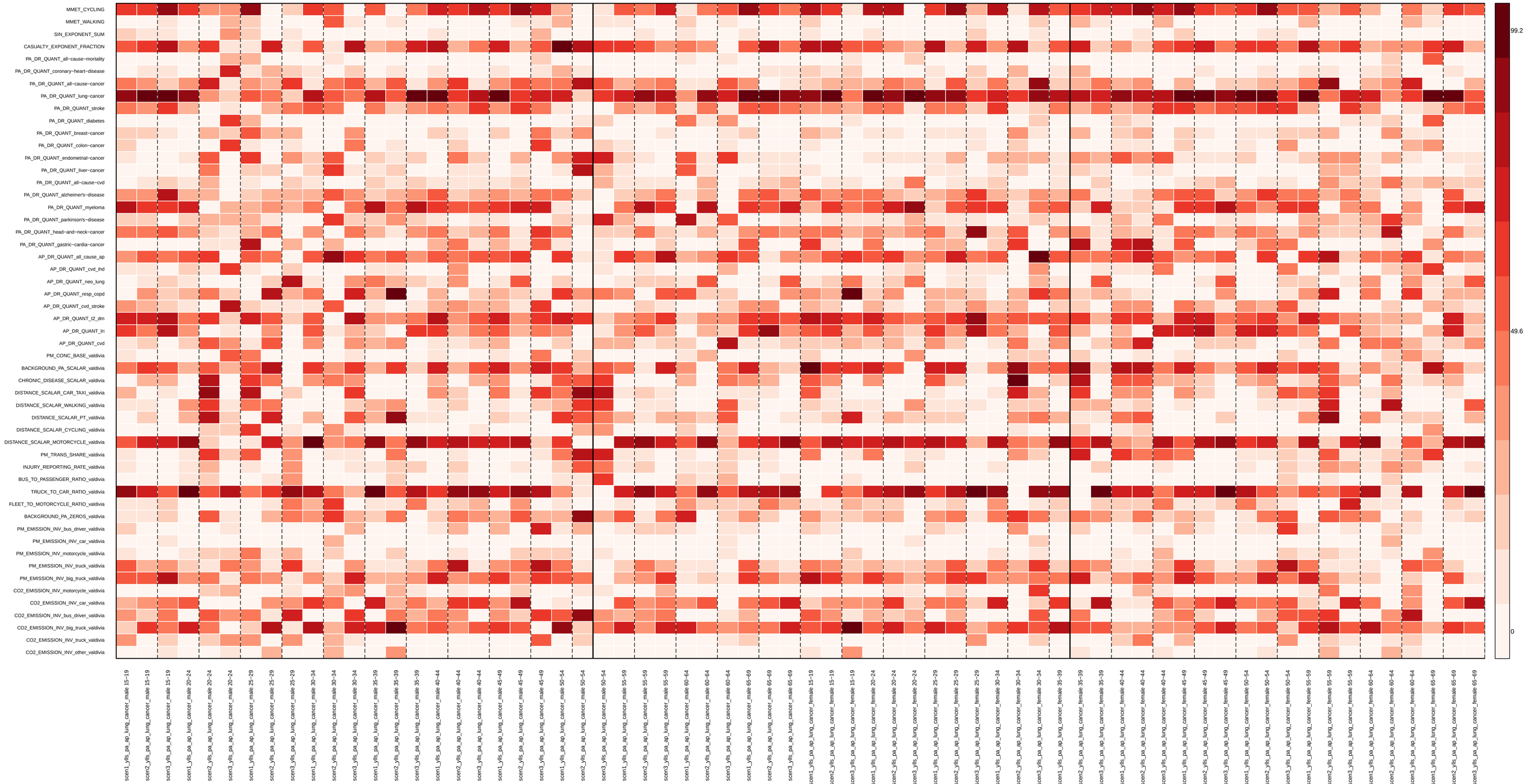
valdivia − pa_ap_IHD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



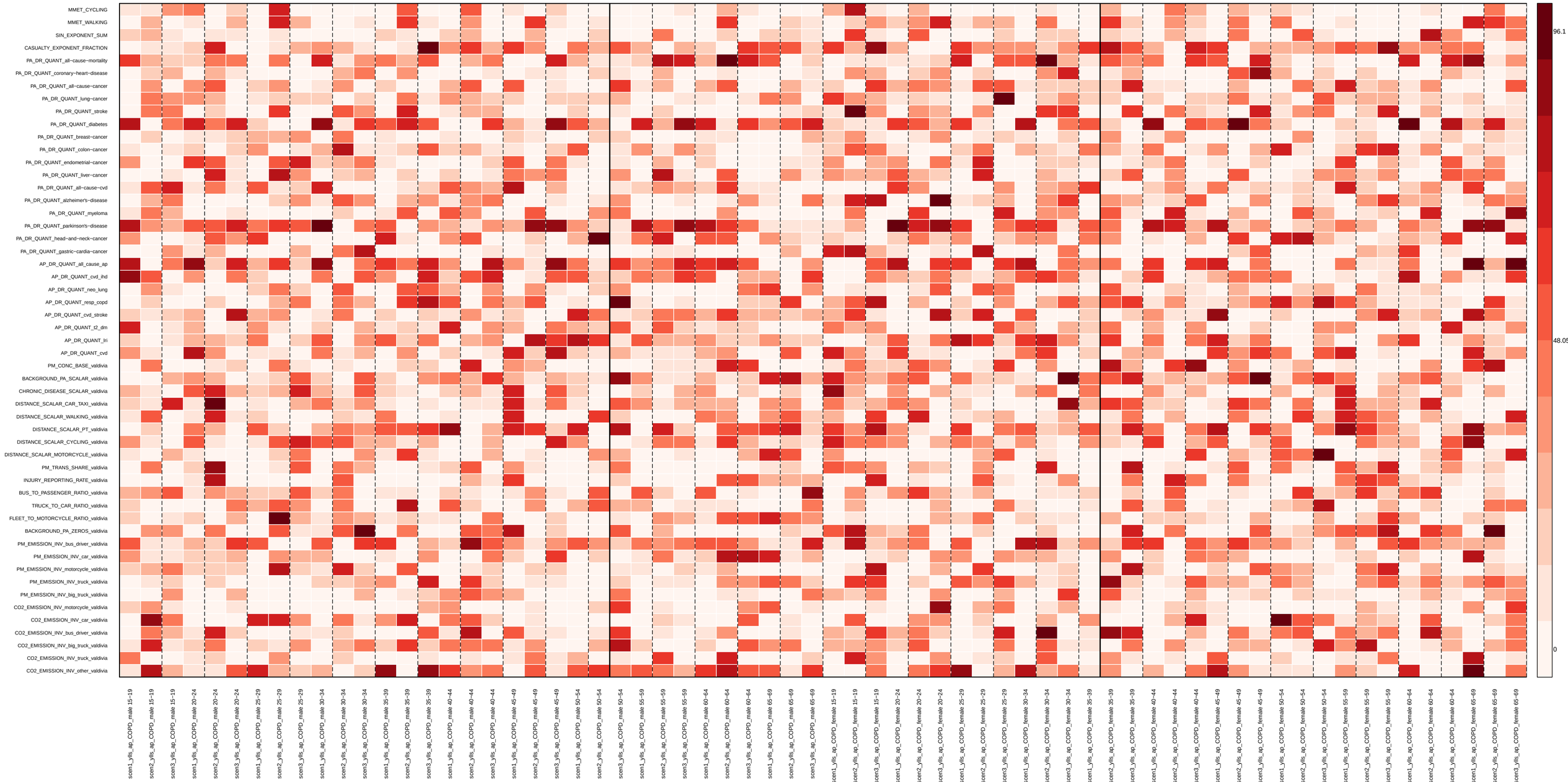
valdivia – pa_total_cancer , No of samples: 10 –
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



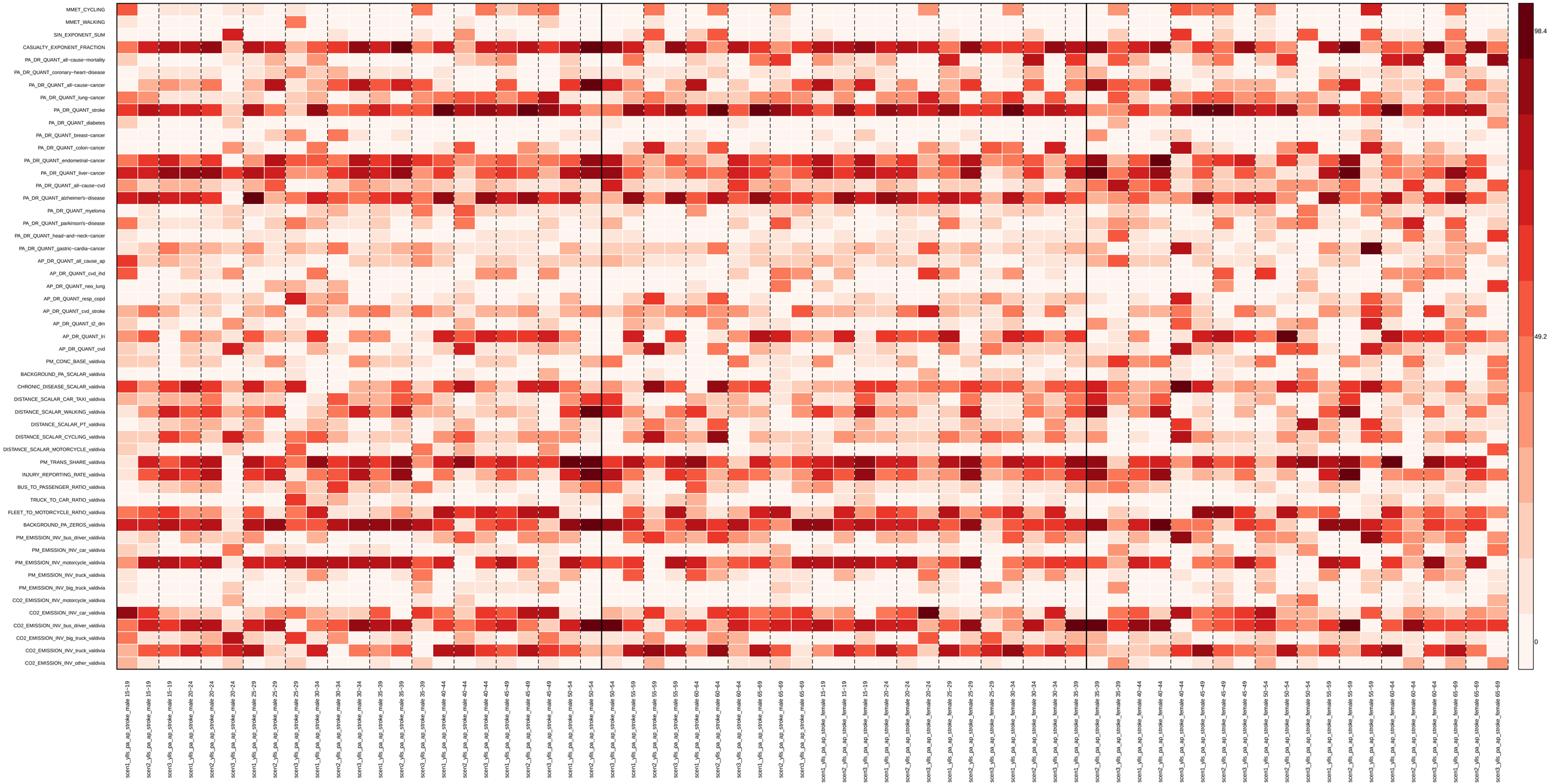
valdivia − pa_ap_lung_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



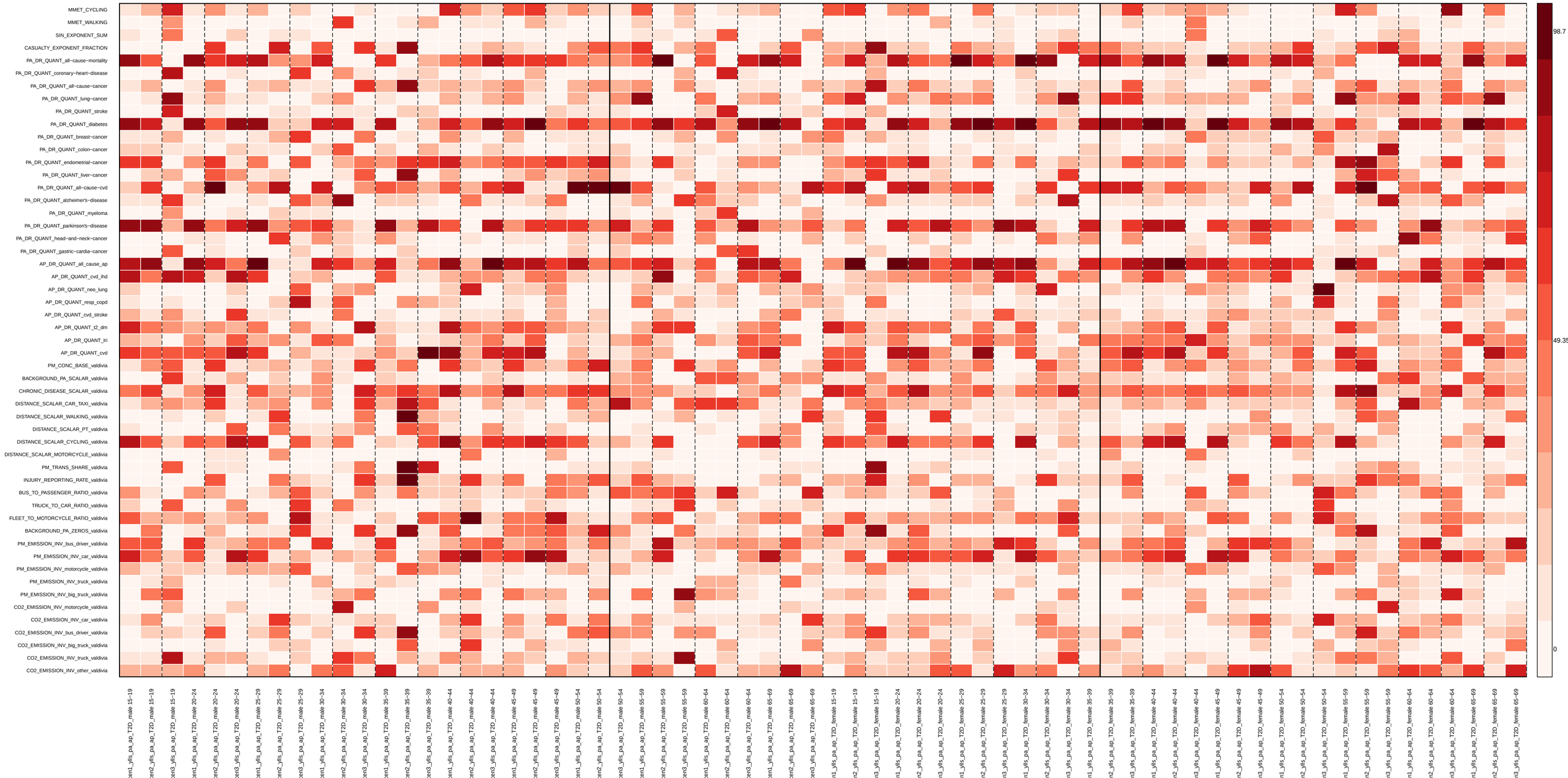
valdivia − ap_COPD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



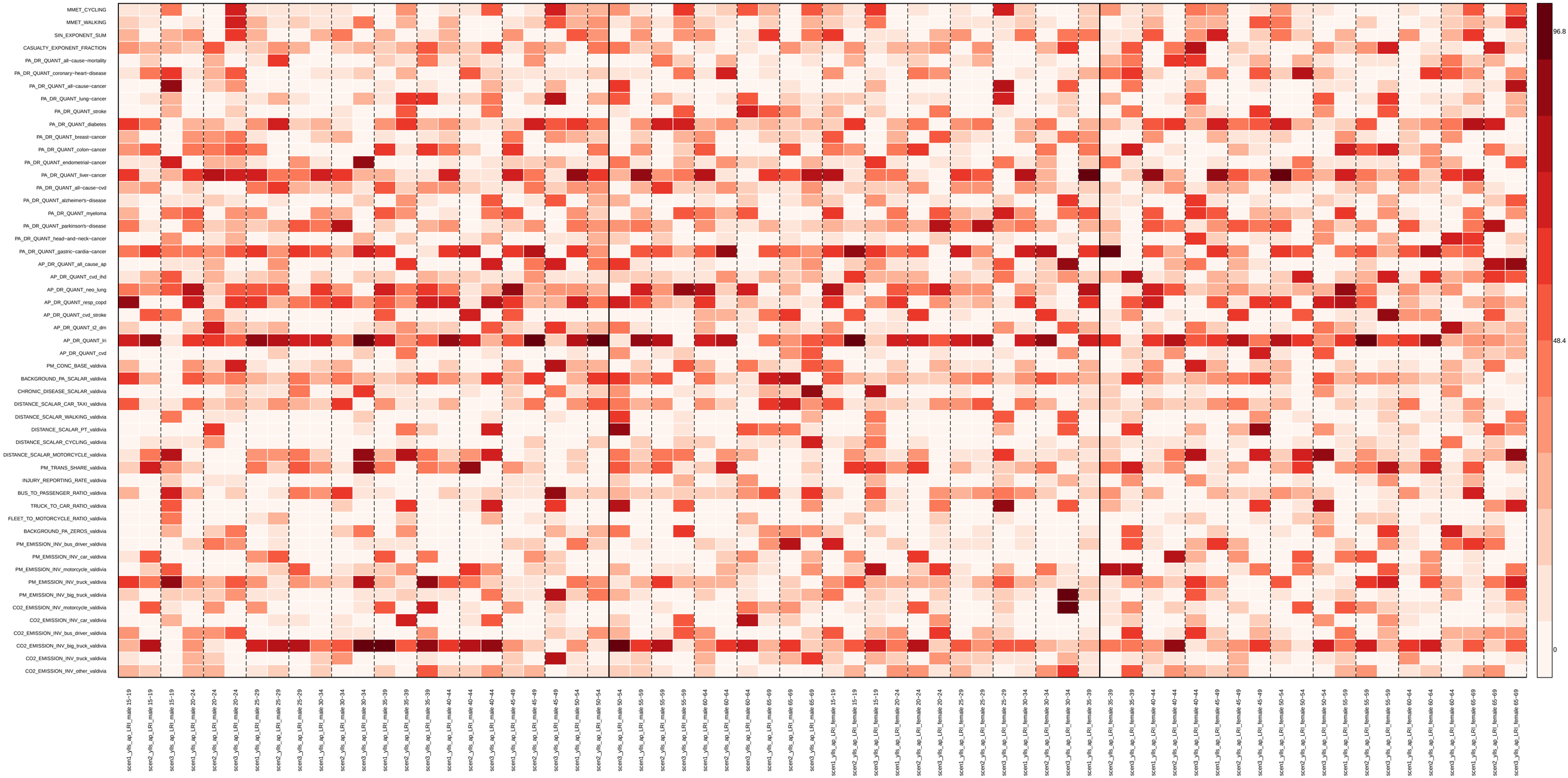
valdivia − pa_ap_stroke , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



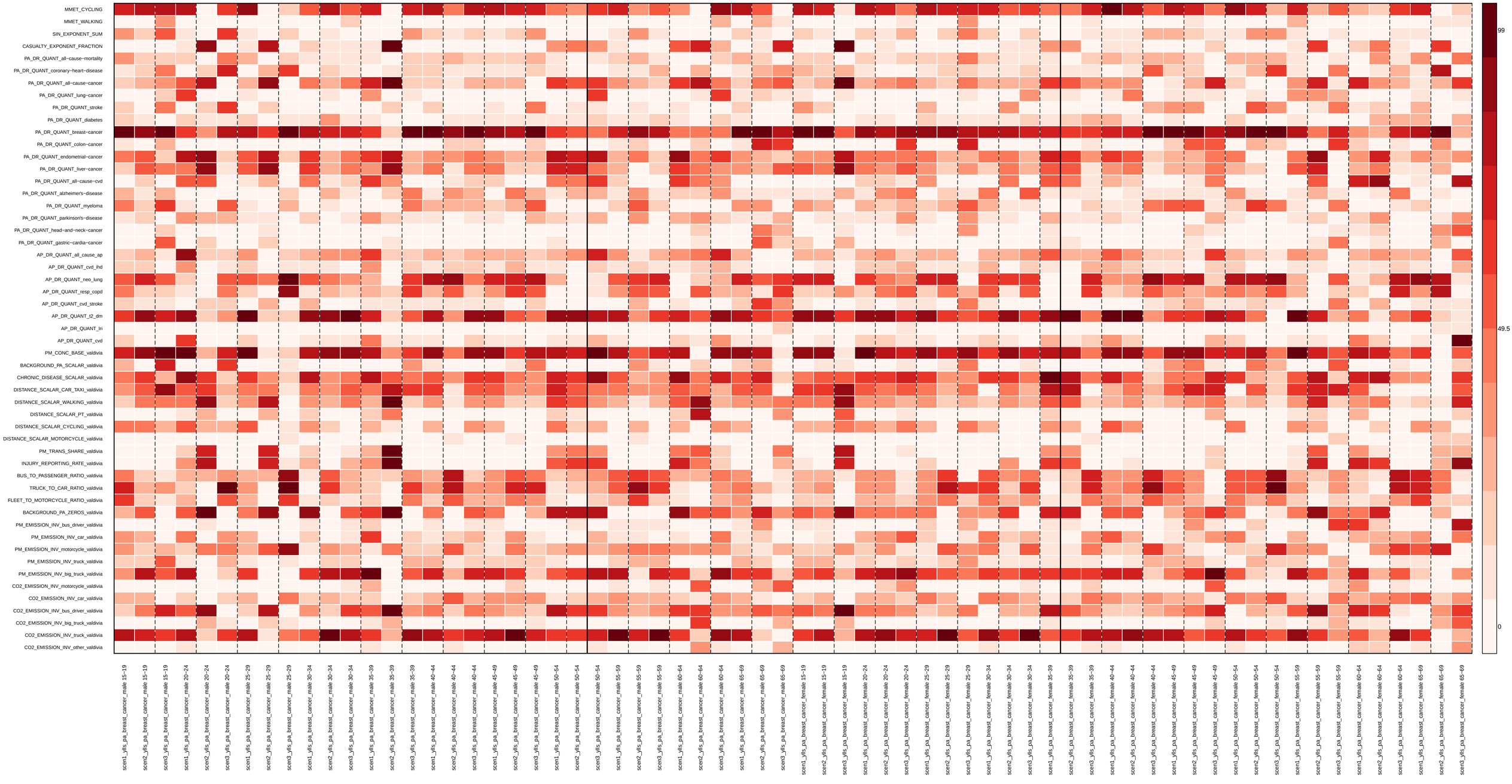
valdivia − pa_ap_T2D , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



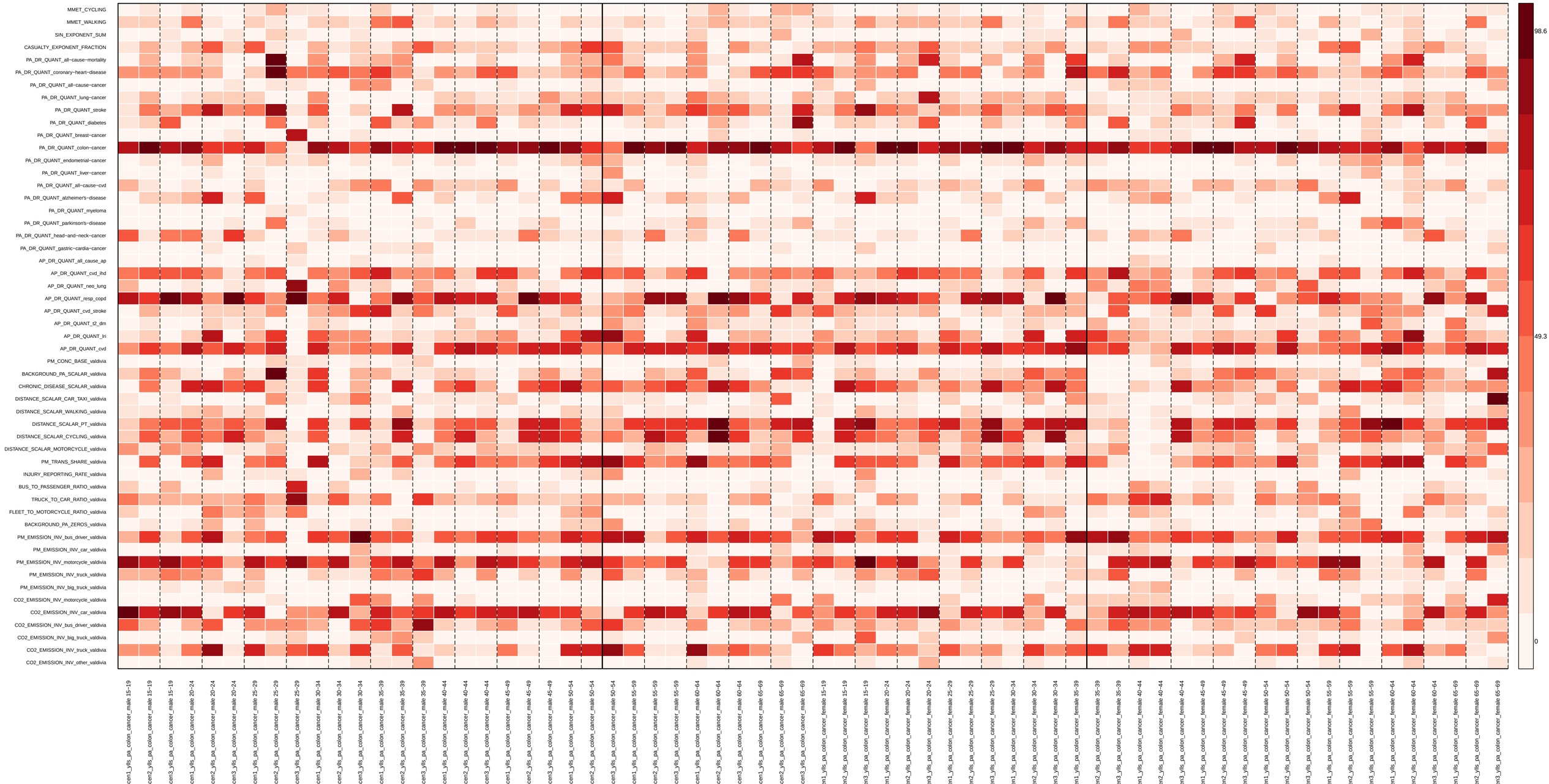
valdivia − ap_LRI , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



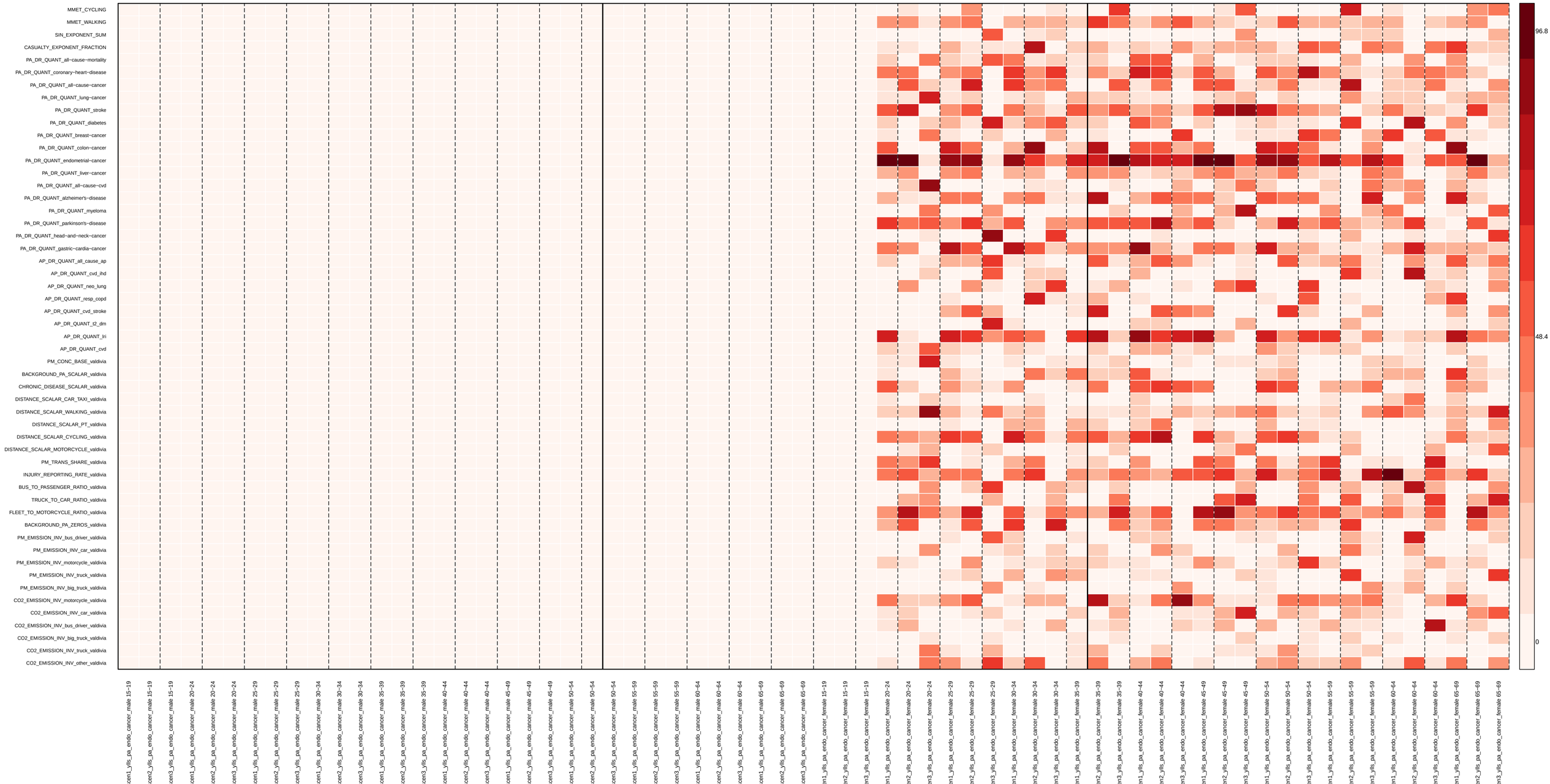
valdivia − pa_breast_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



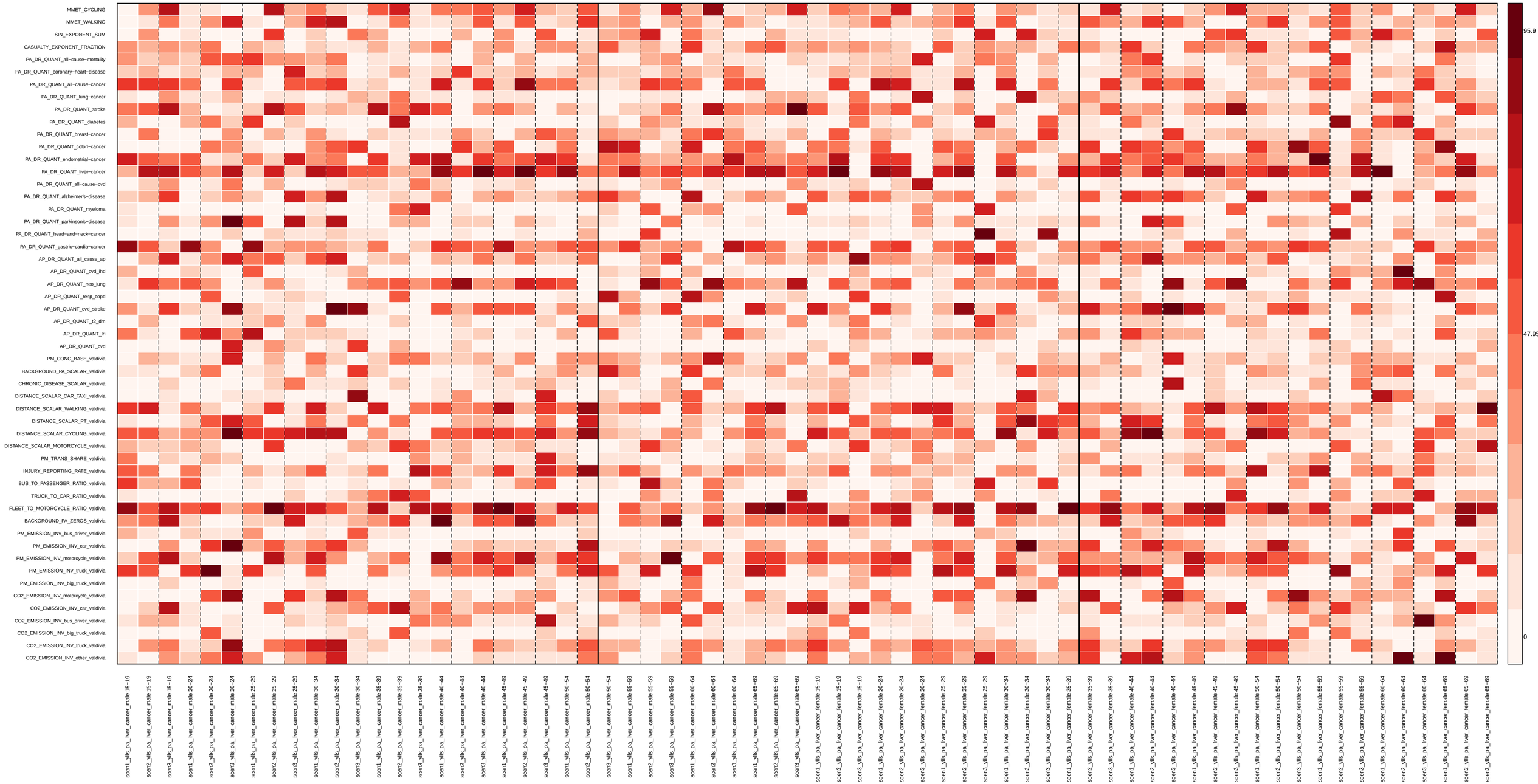
valdivia − pa_colon_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



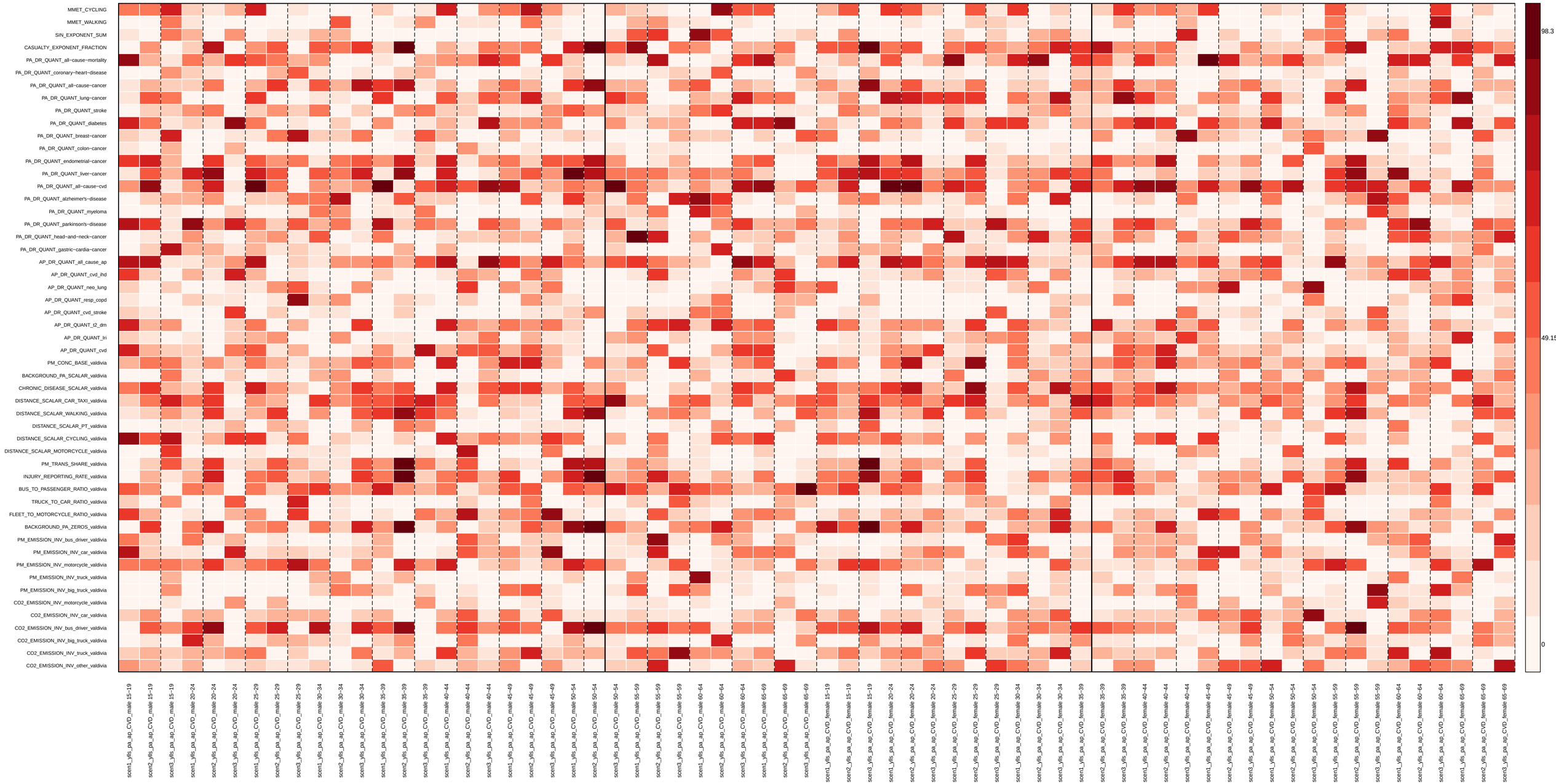
valdivia − pa_endo_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



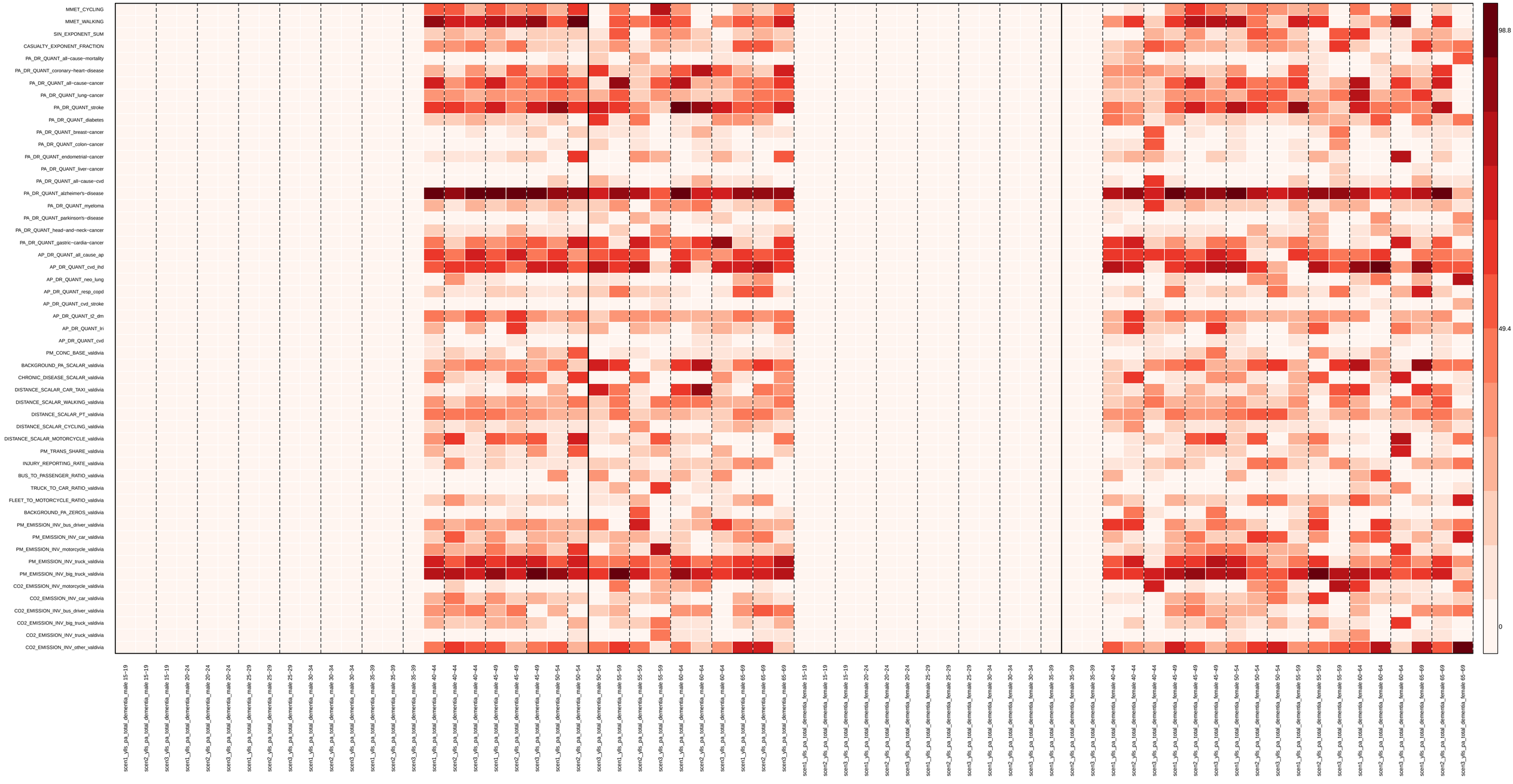
valdivia − pa_liver_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



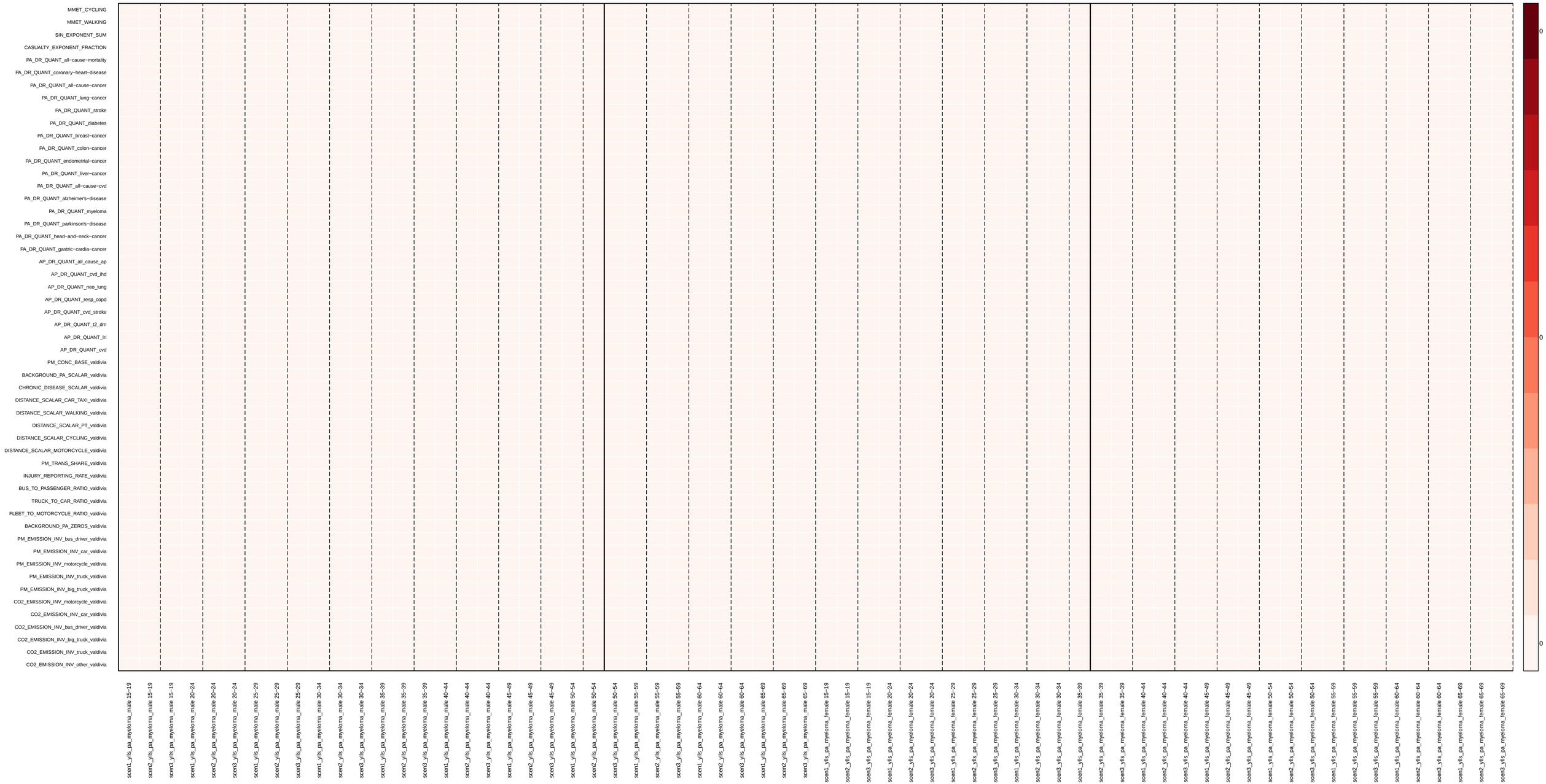
valdivia − pa_ap_CVD , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



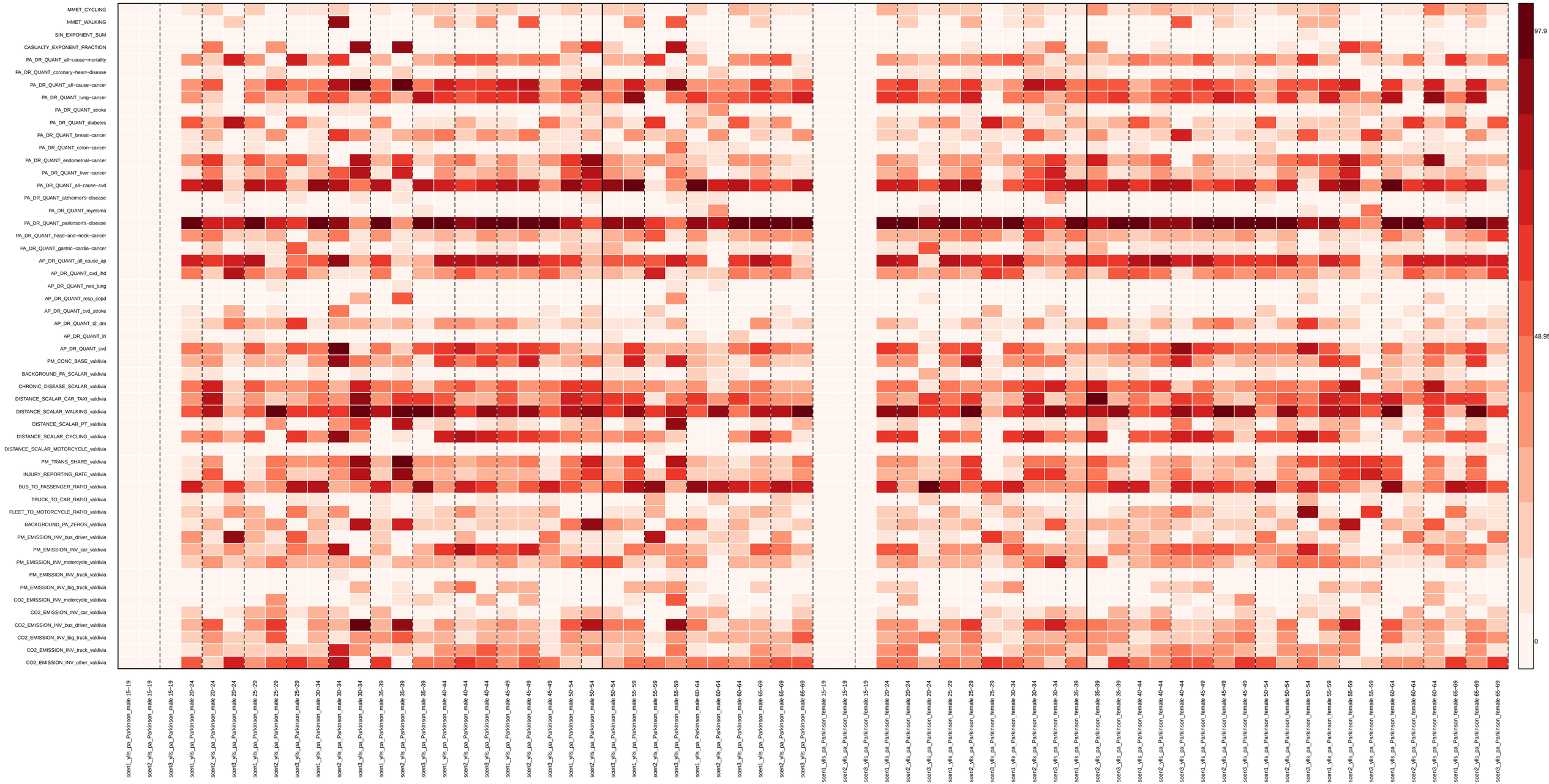
valdivia − pa_total_dementia , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



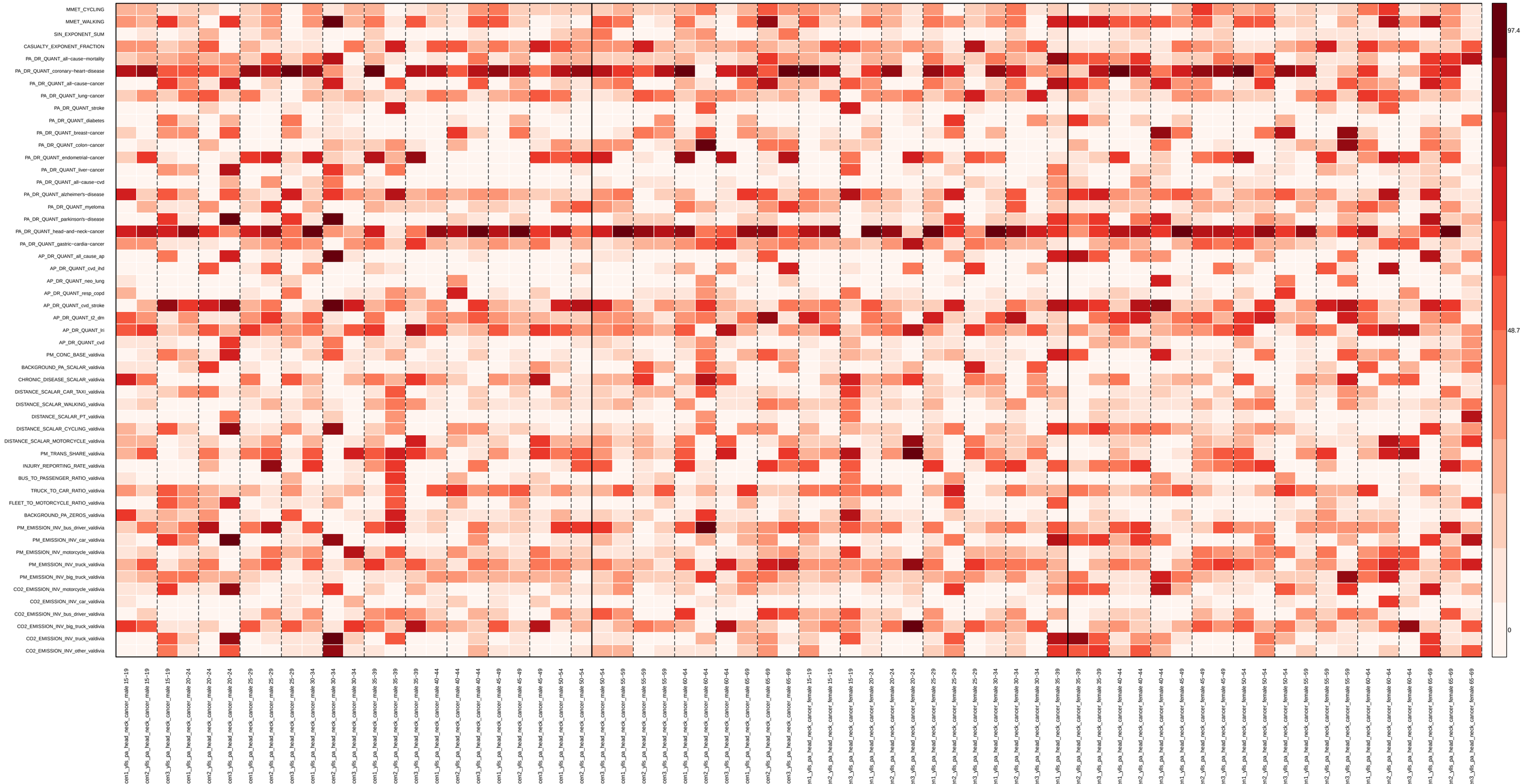
valdivia − pa_myeloma , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



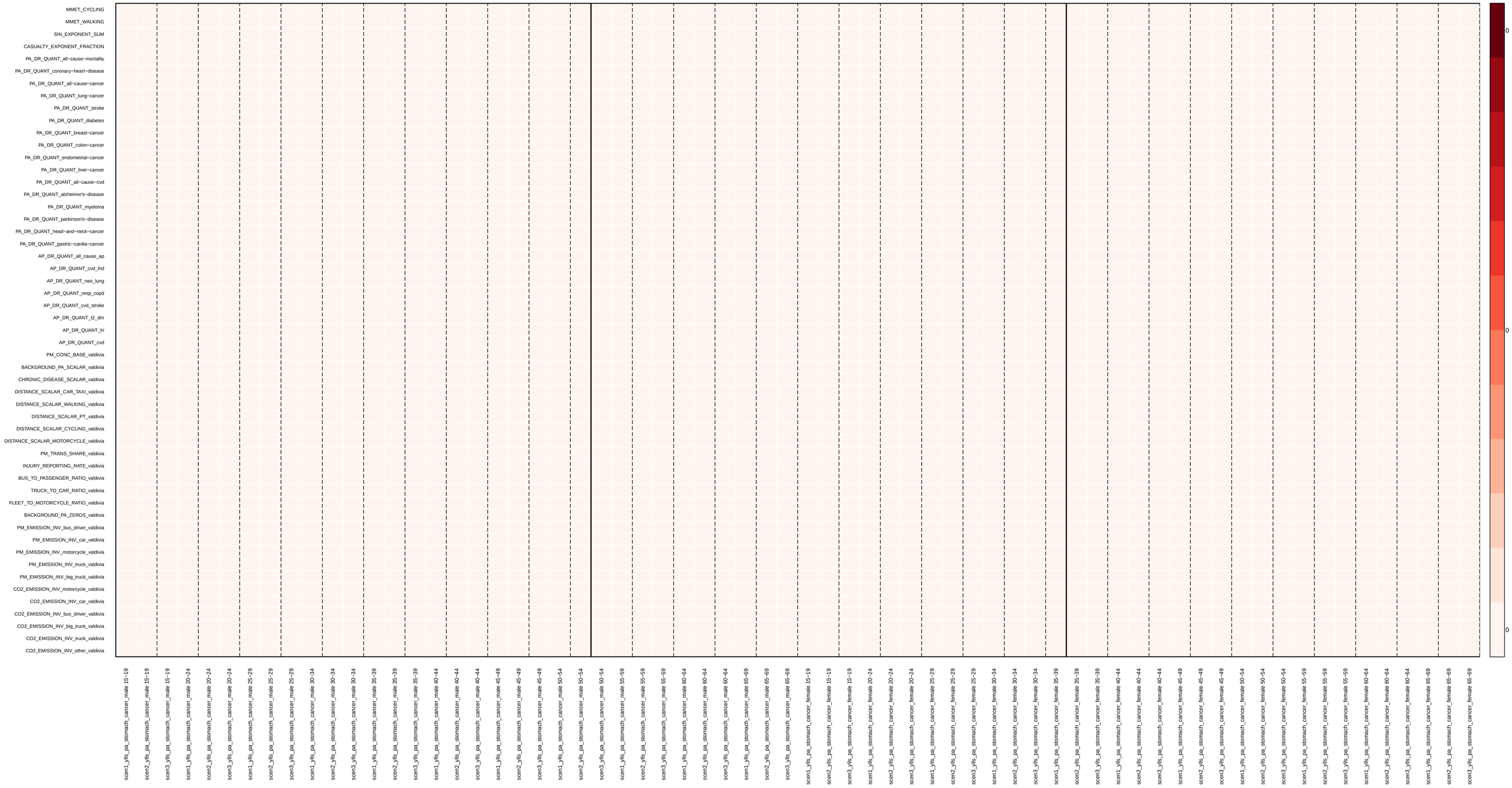
valdivia − pa_Parkinson , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



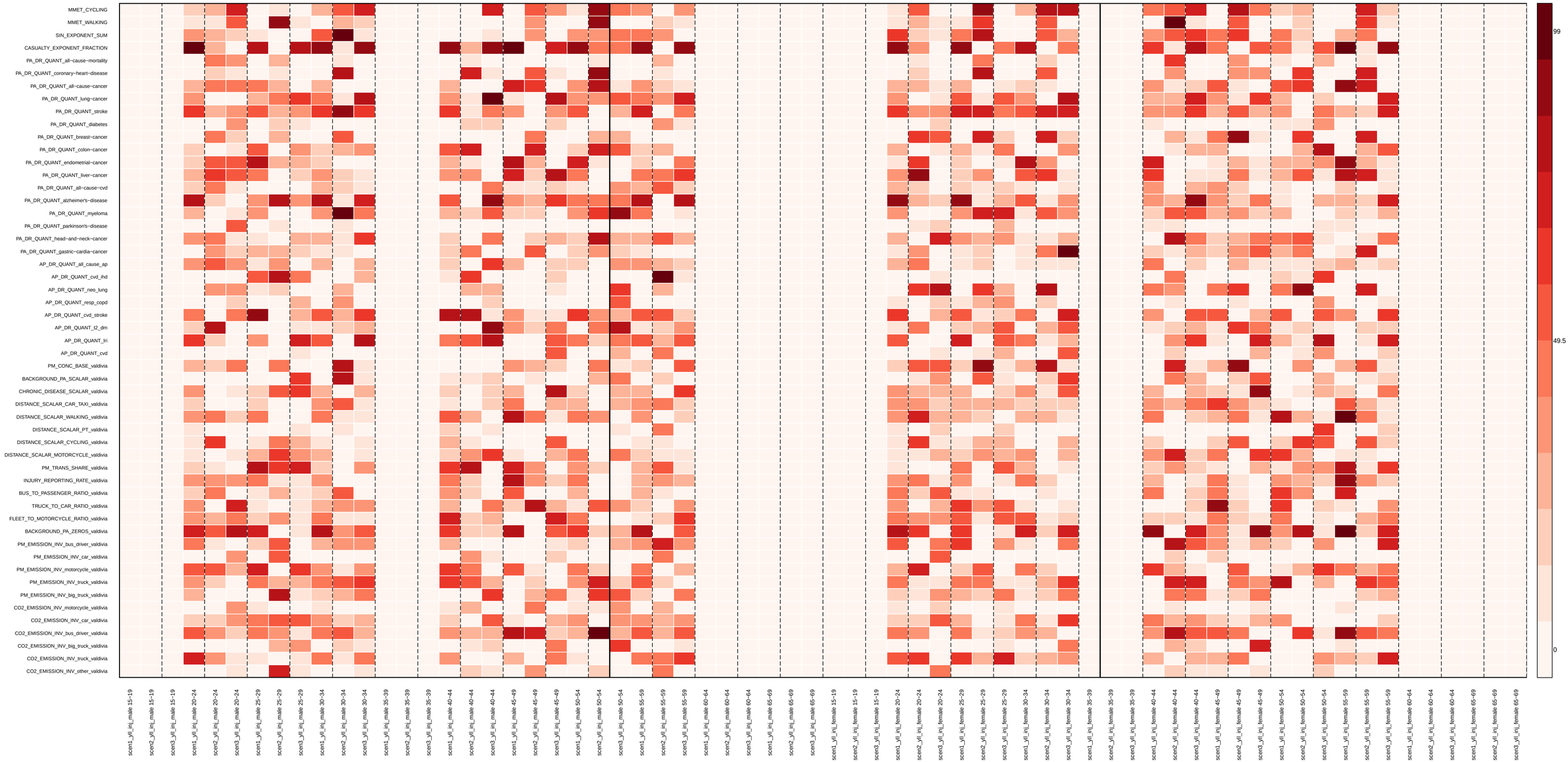
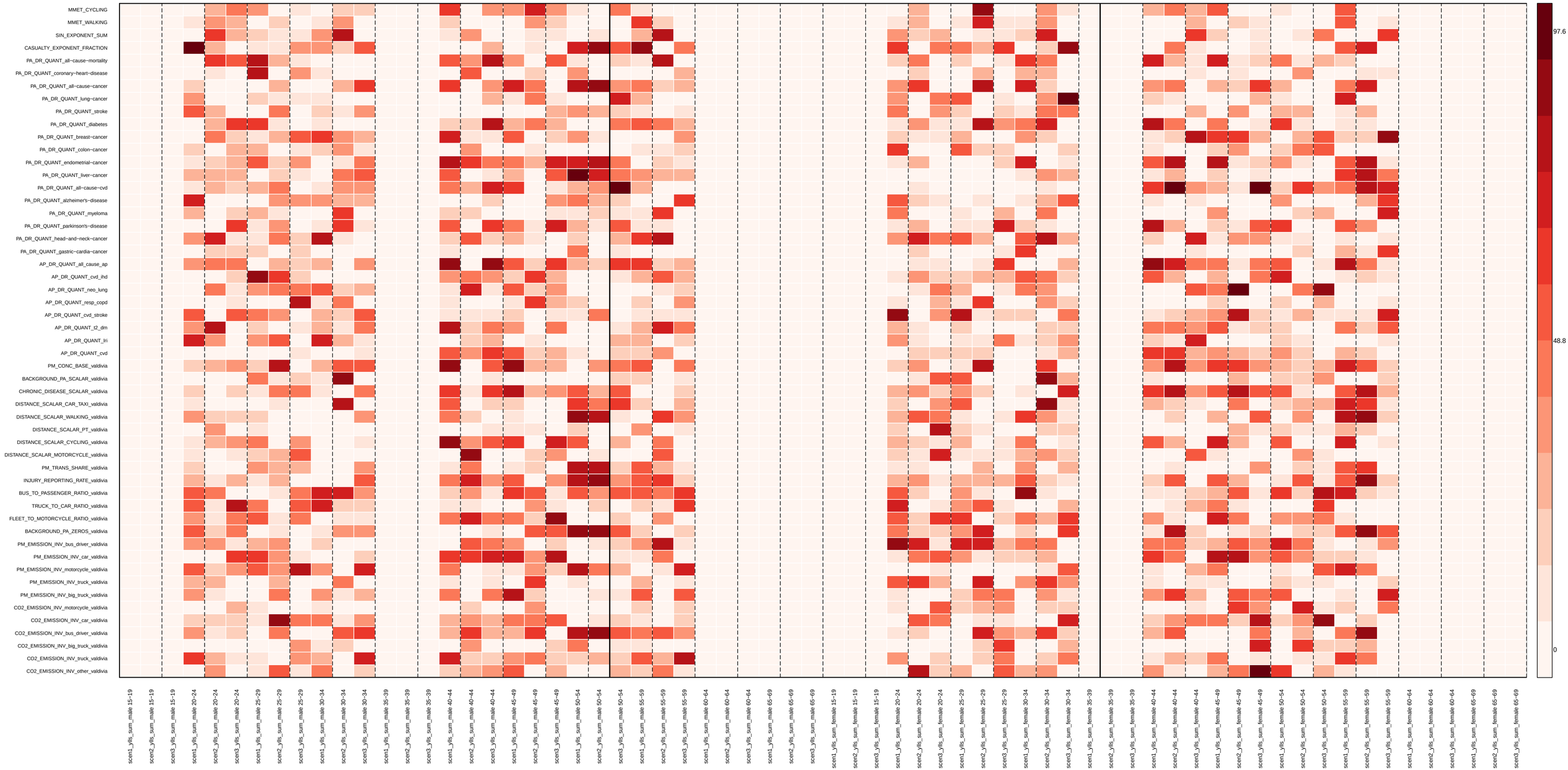
valdivia − pa_head_neck_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



valdivia − pa_stomach_cancer , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



valdivia − inj , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?



valdivia − sum , No of samples: 10 −
By how much (%) could we reduce uncertainty in the outcome if we knew this parameter perfectly?