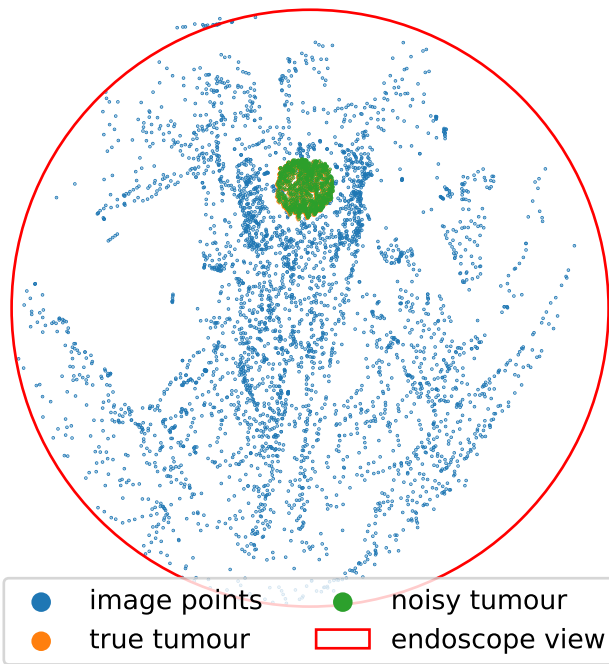
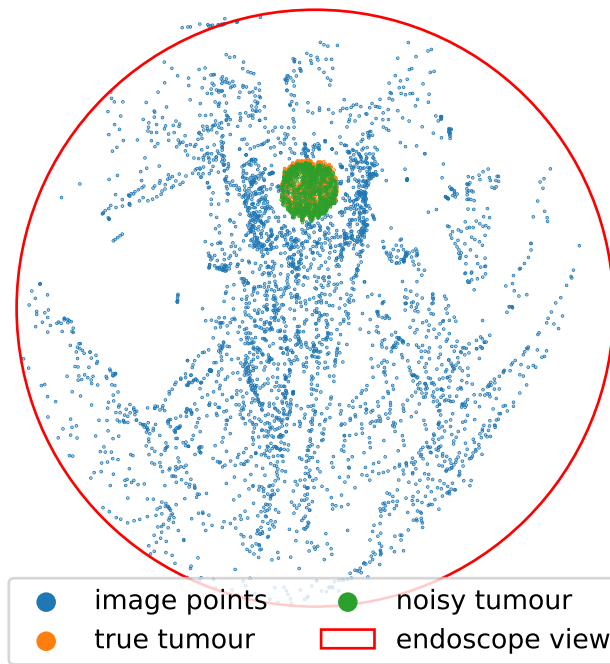


visualising tumour misalignment at 0.12 error



visualising tumour misalignment at 0.25 error



visualising tumour misalignment at 0.5 error

