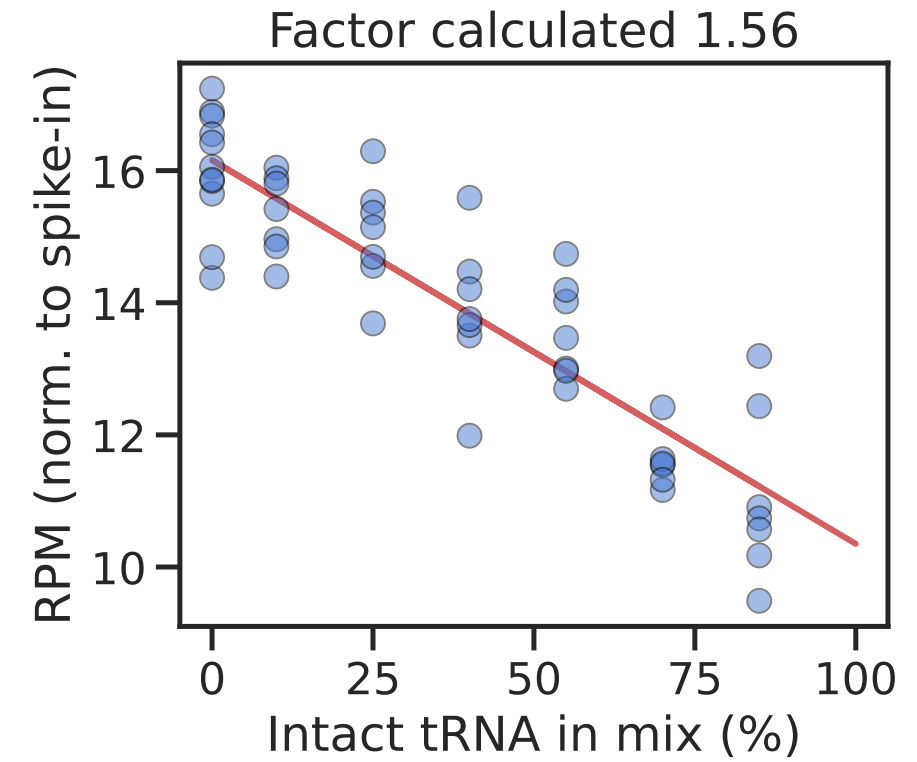
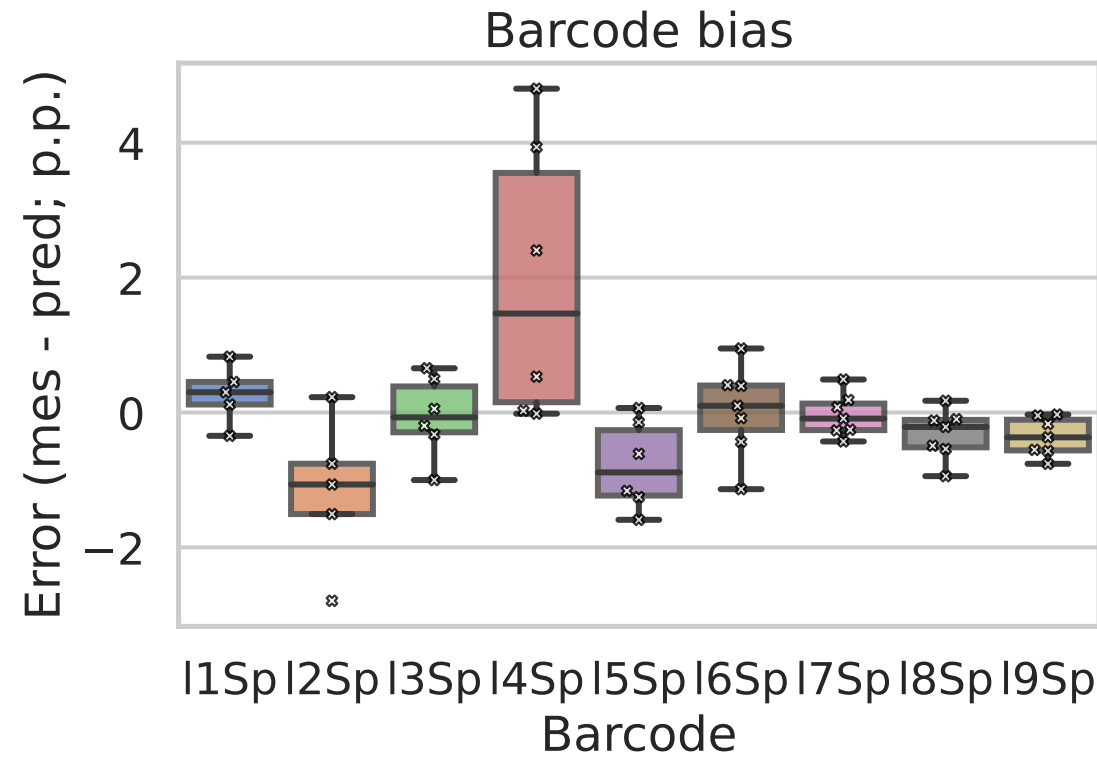
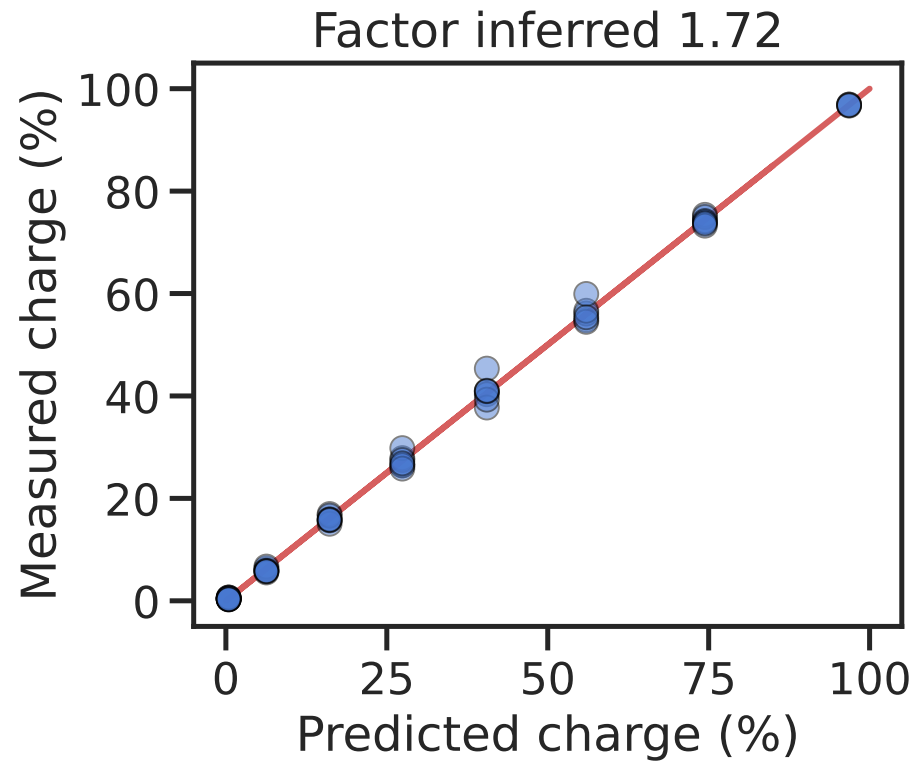
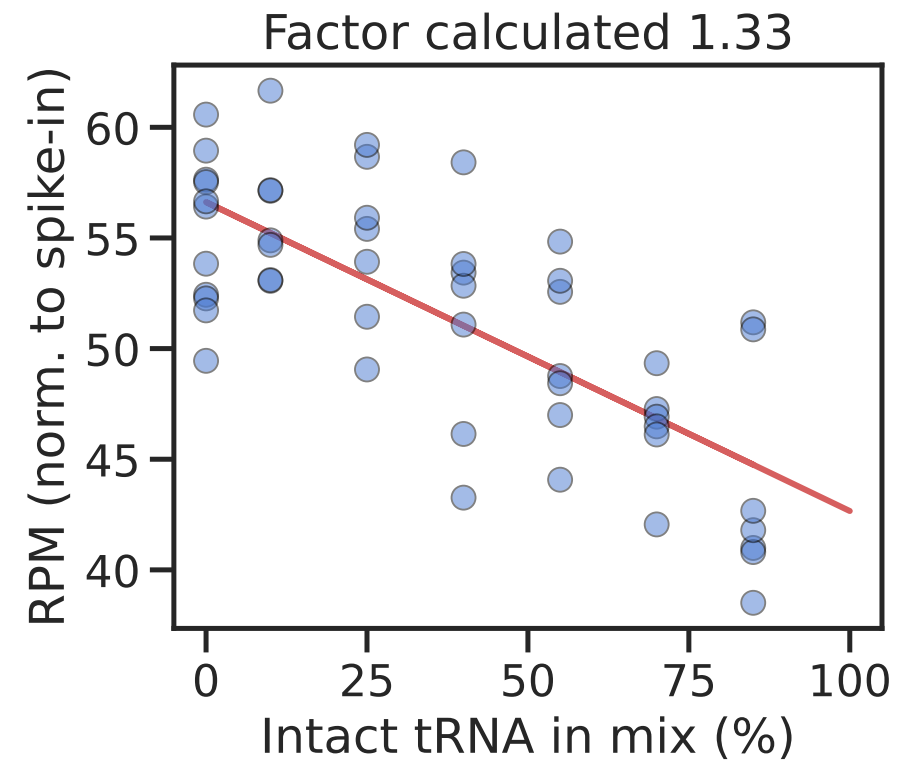
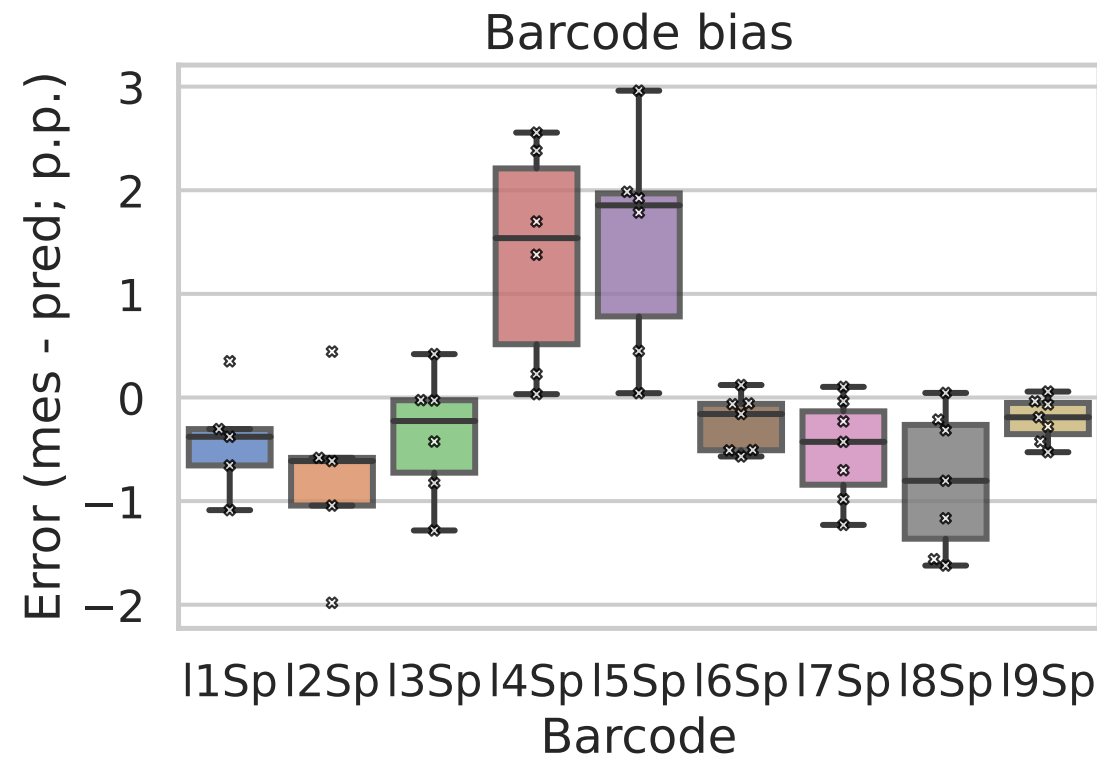
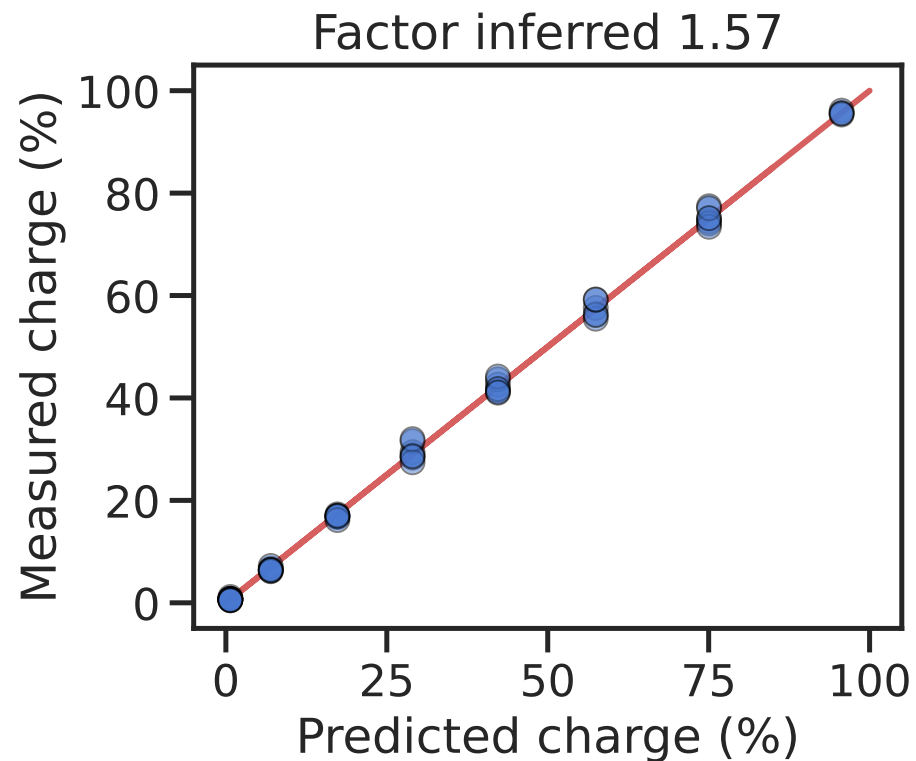


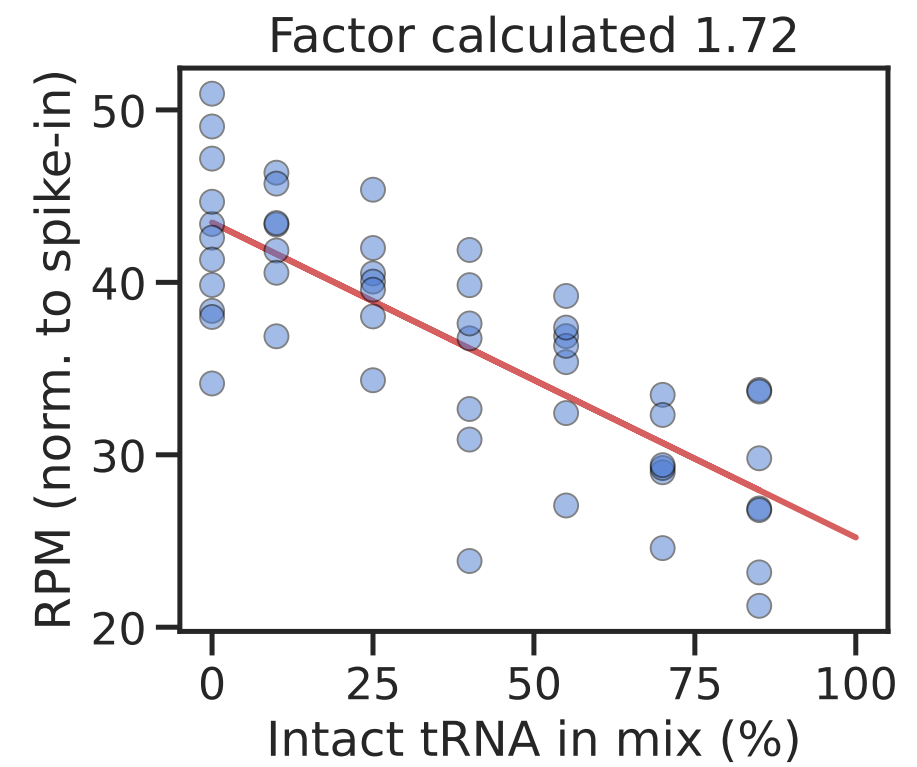
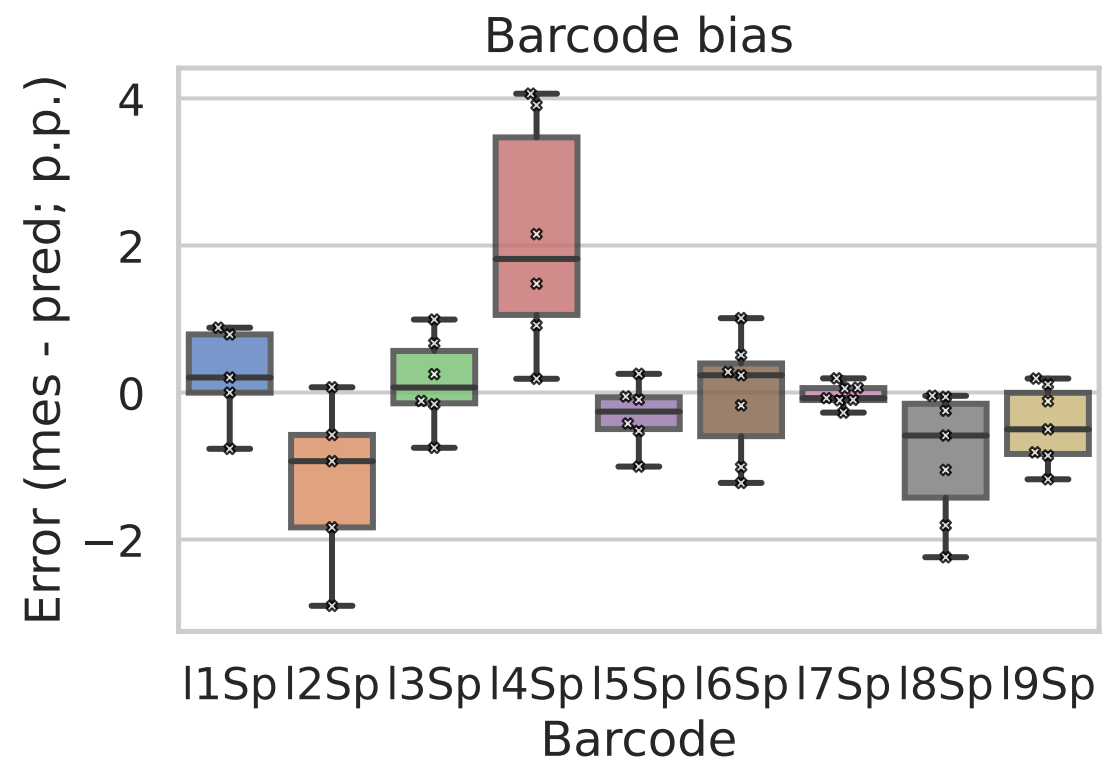
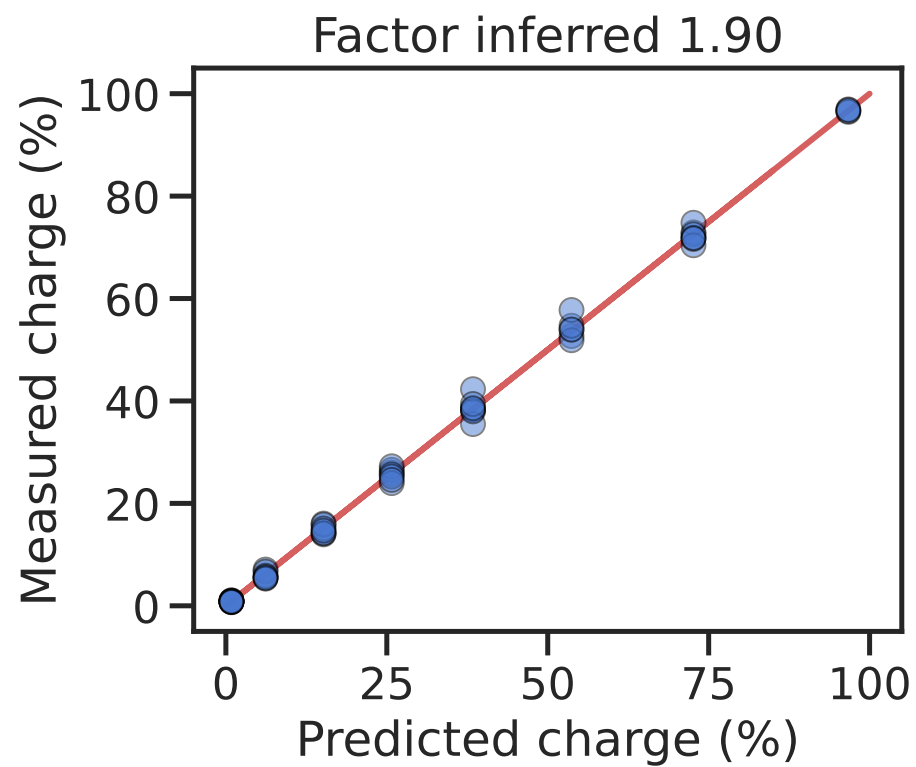
Ala-TGC-4-1



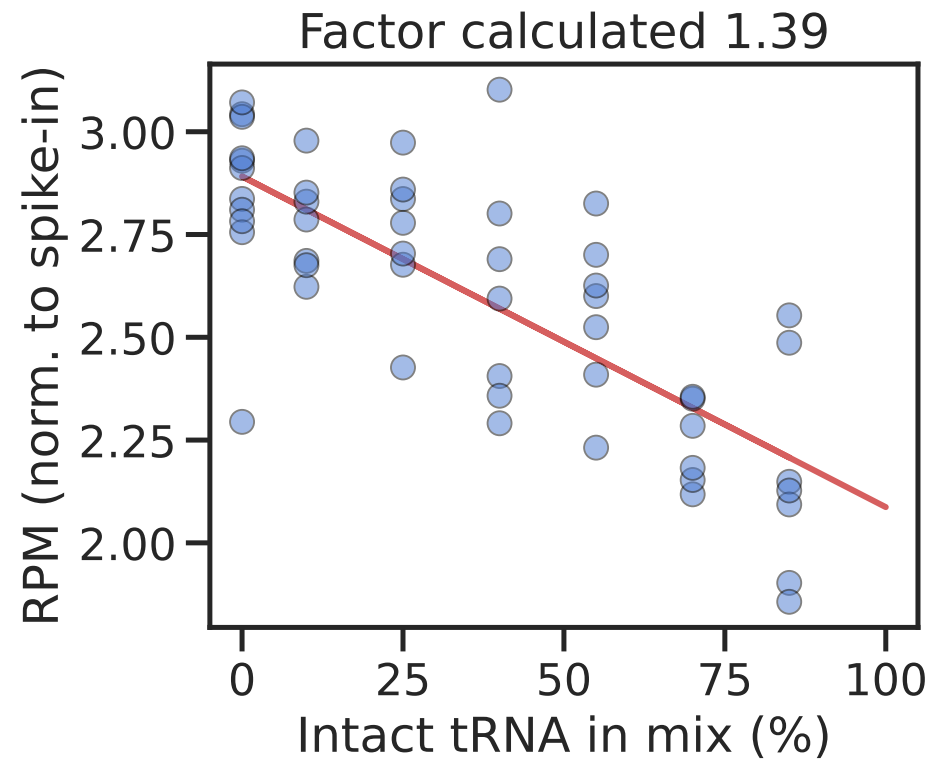
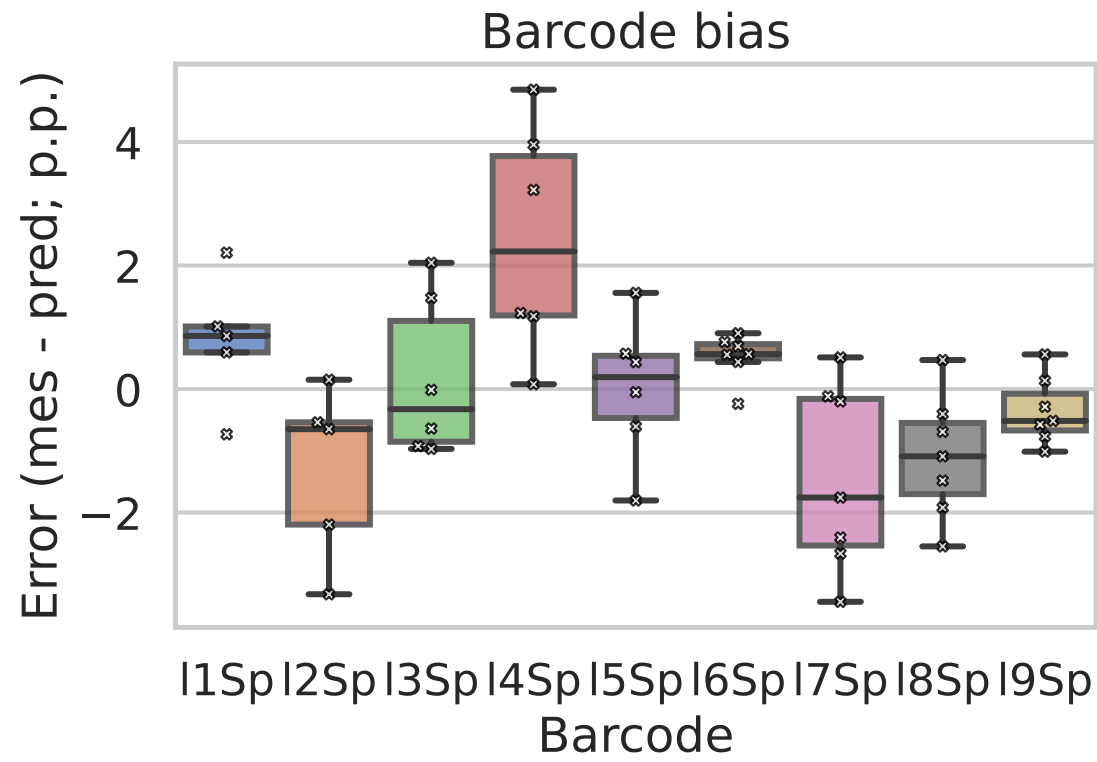
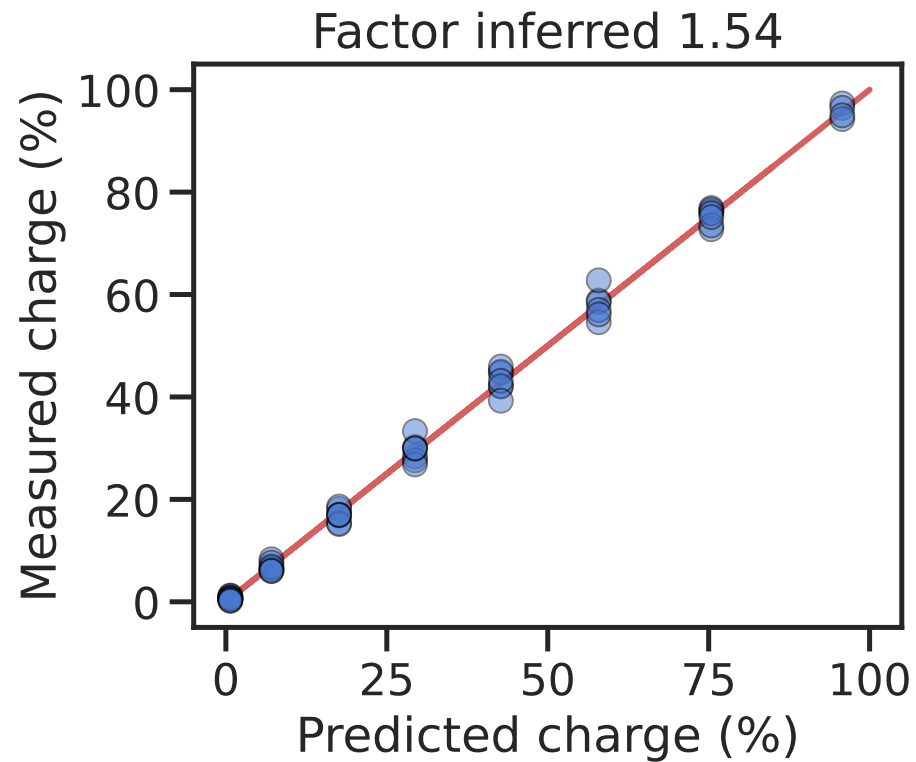
Lys-TTT-3-1



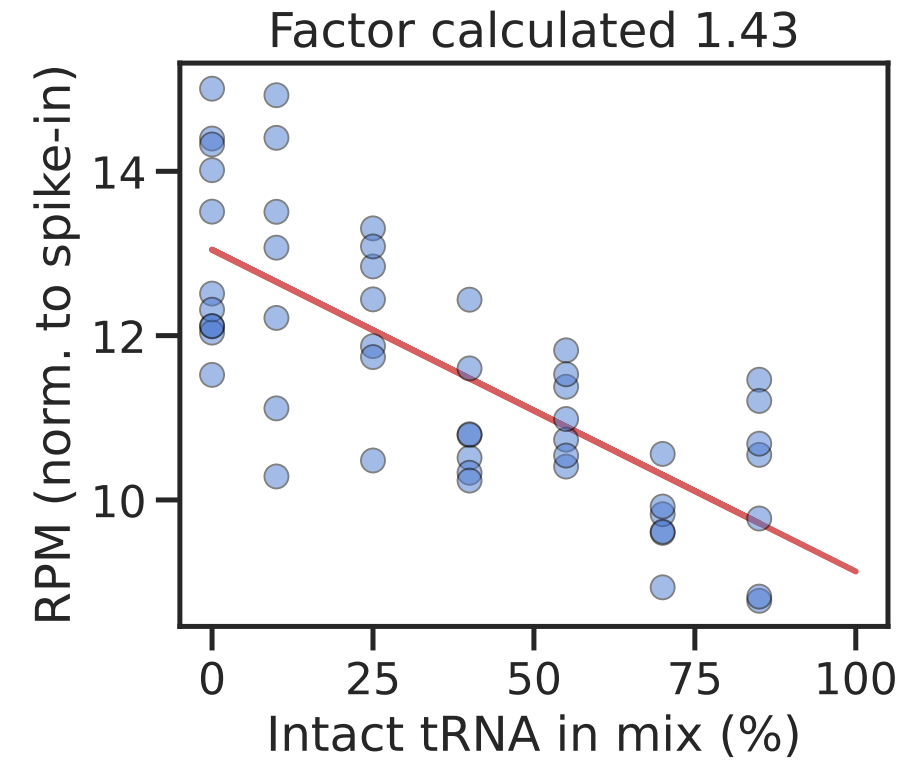
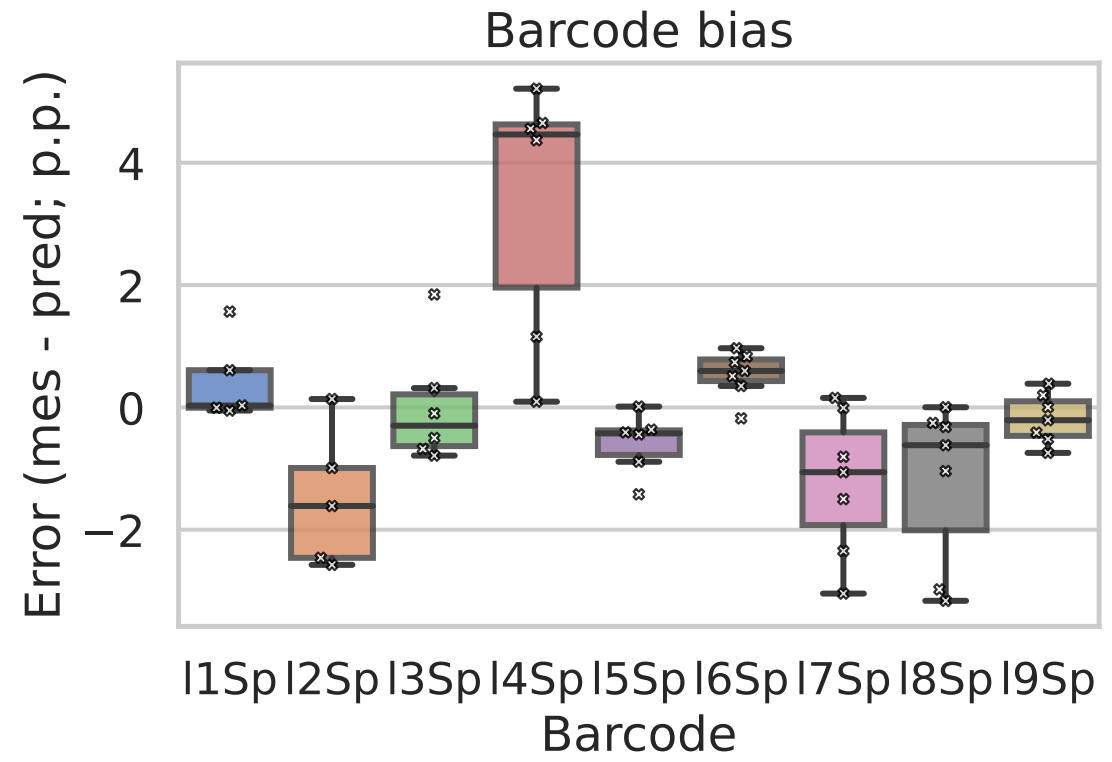
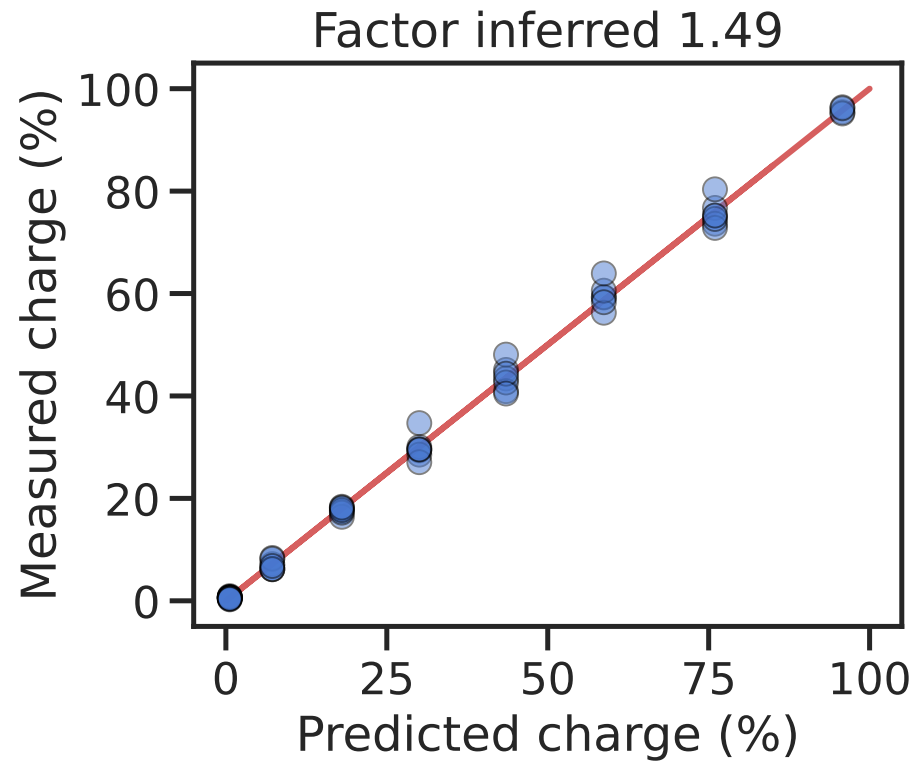
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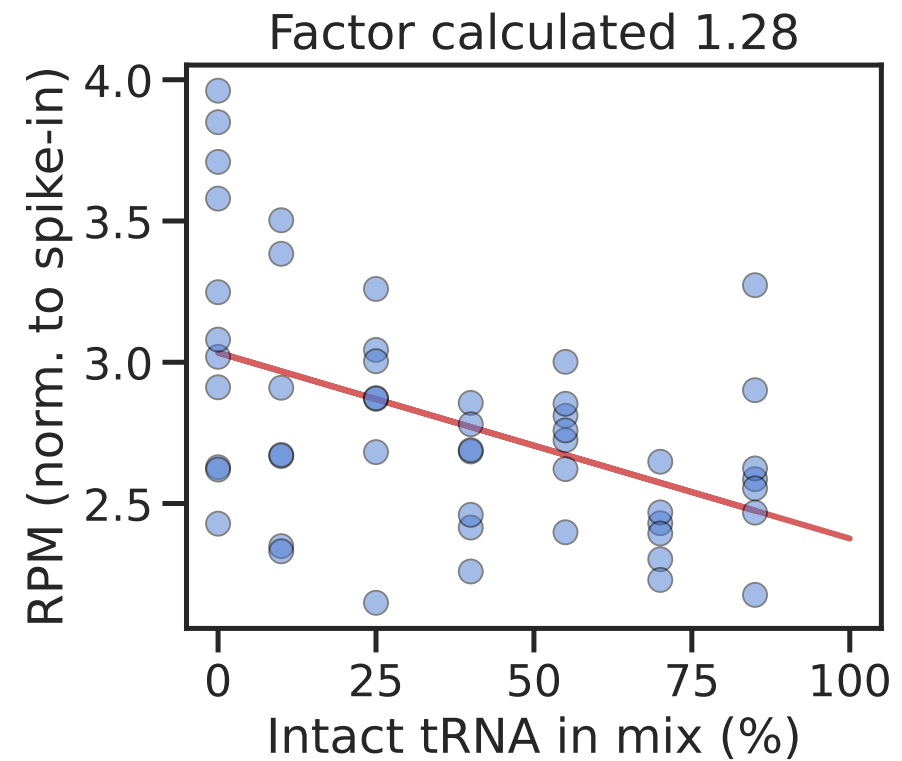
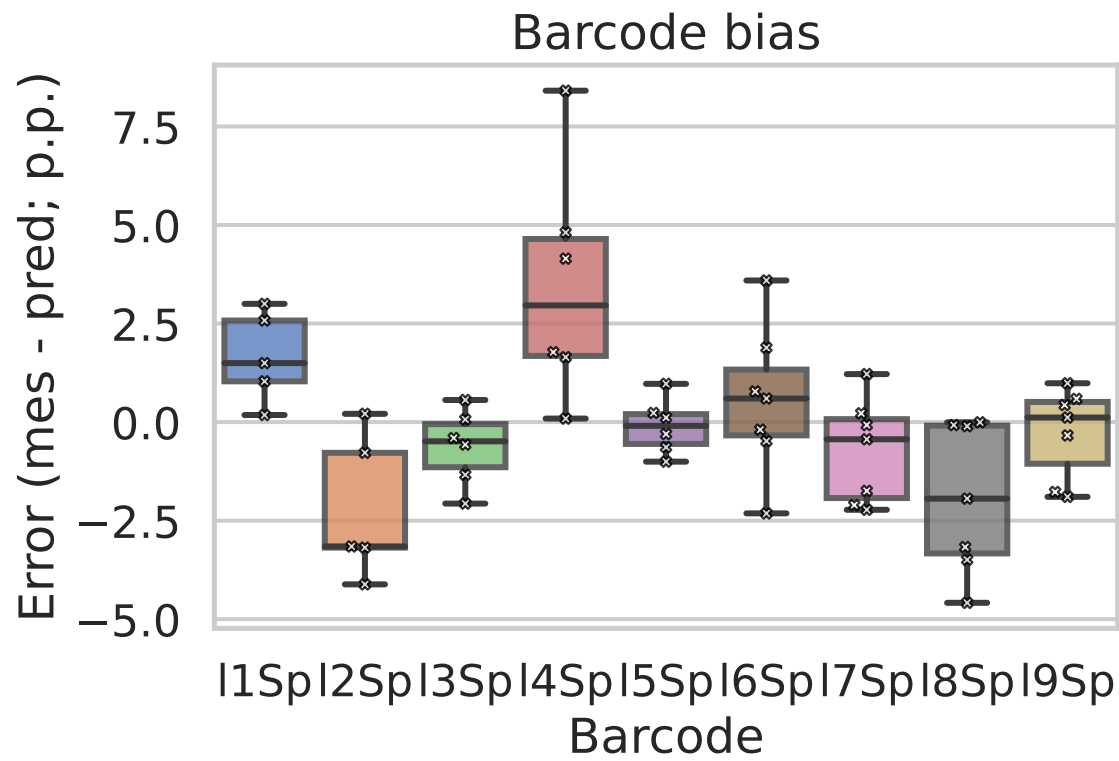
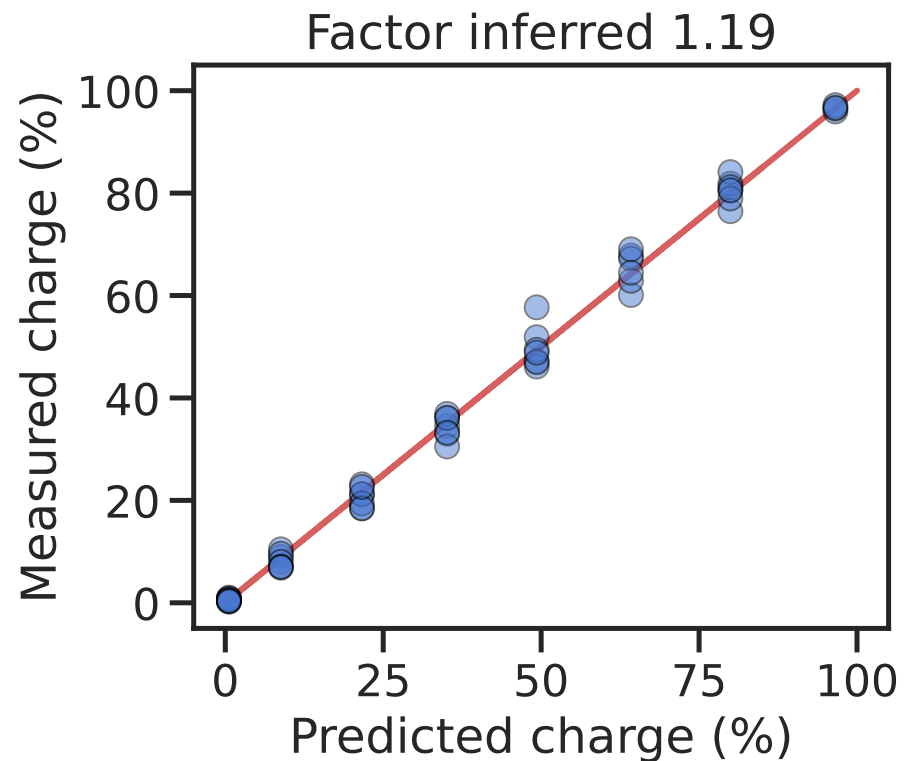
Ala-AGC-5-1



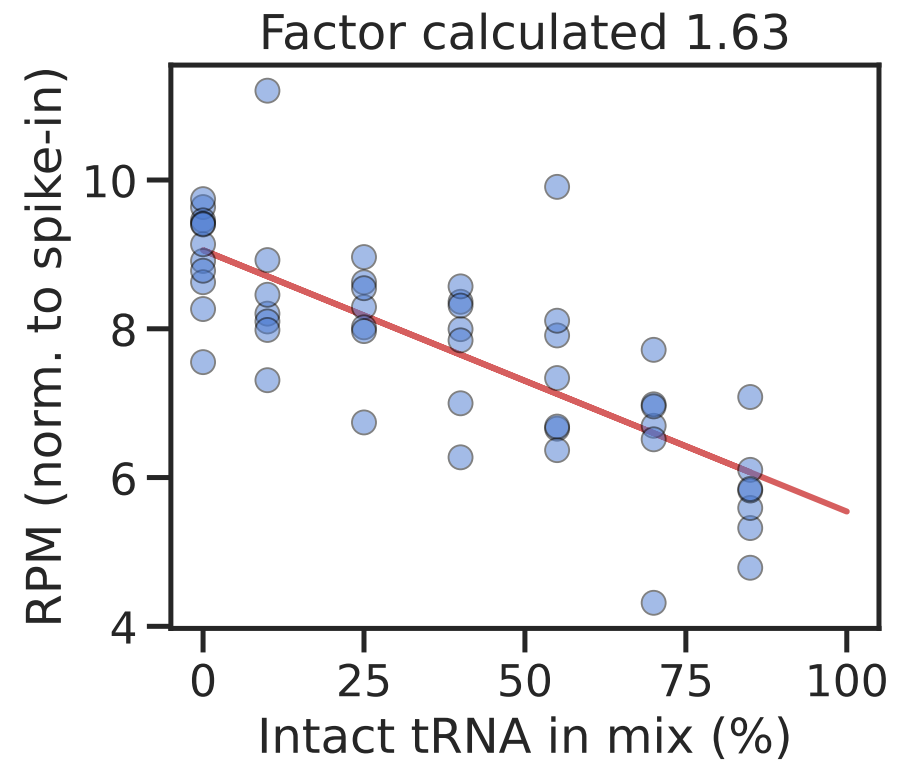
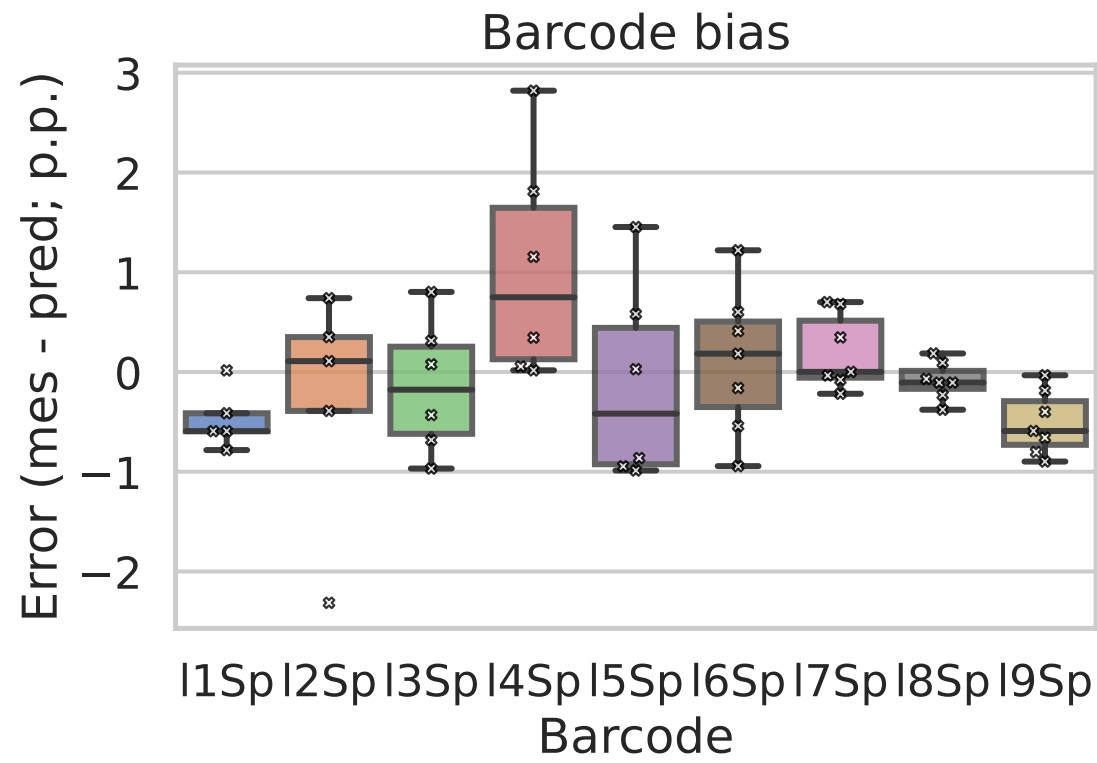
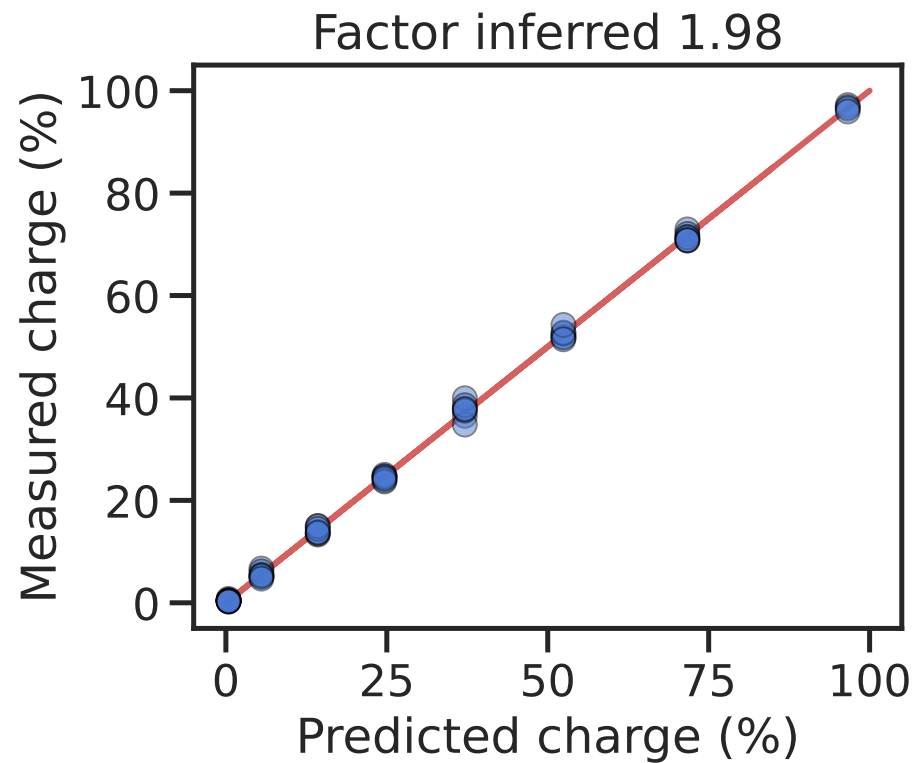
Thr-AGT-1-1



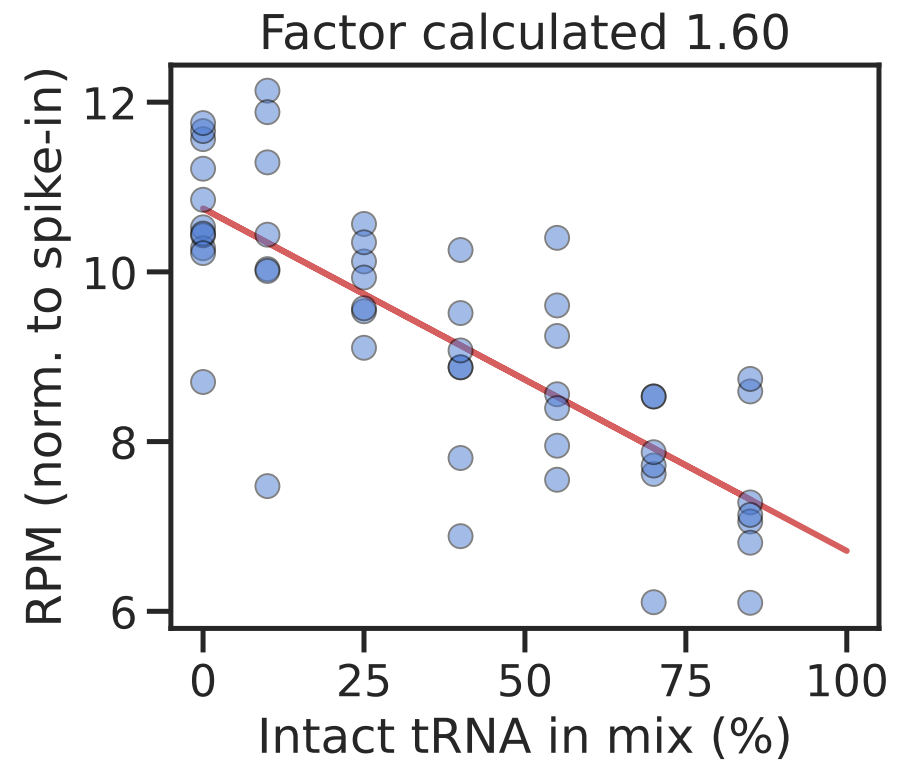
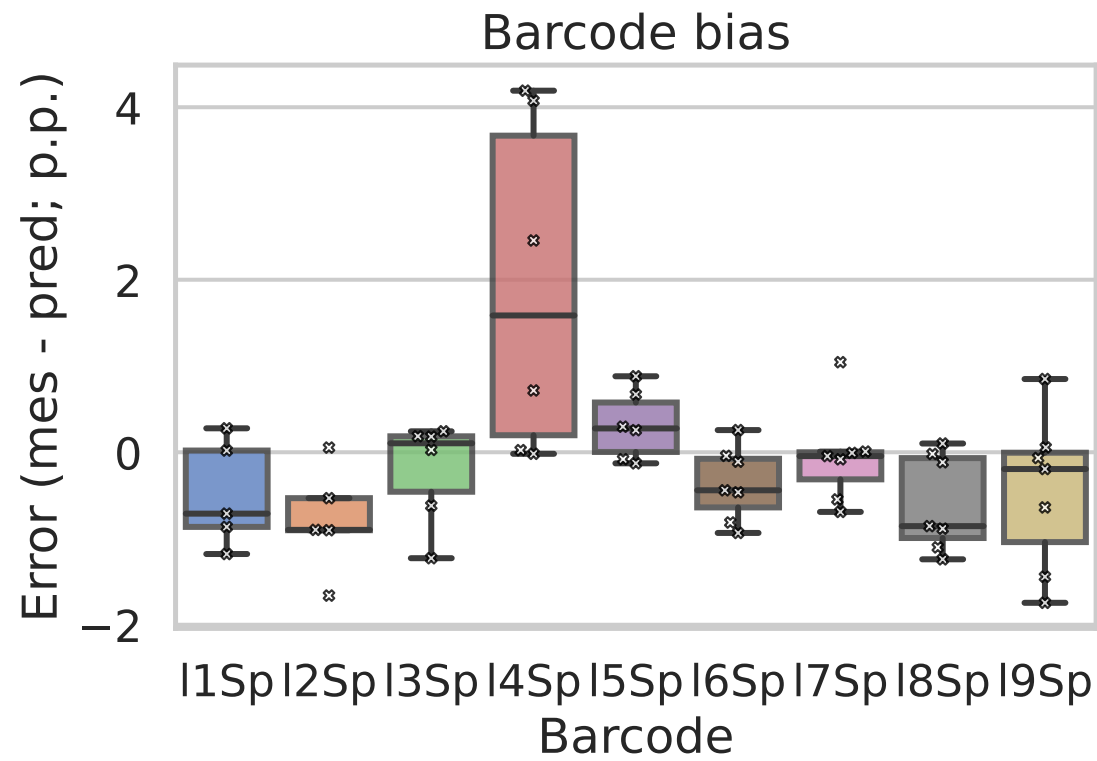
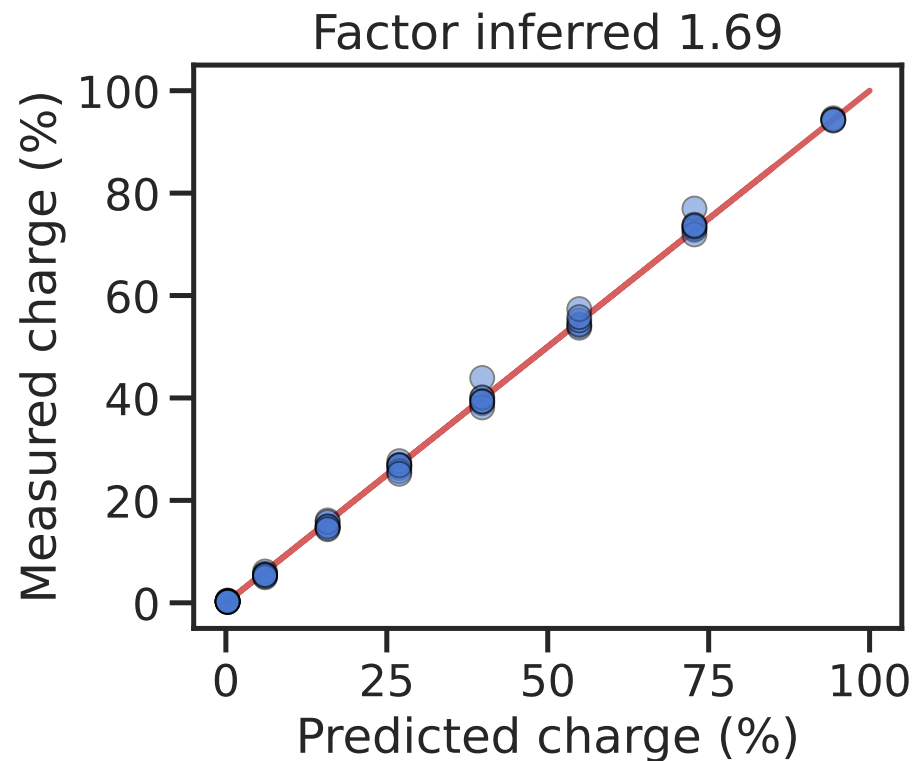
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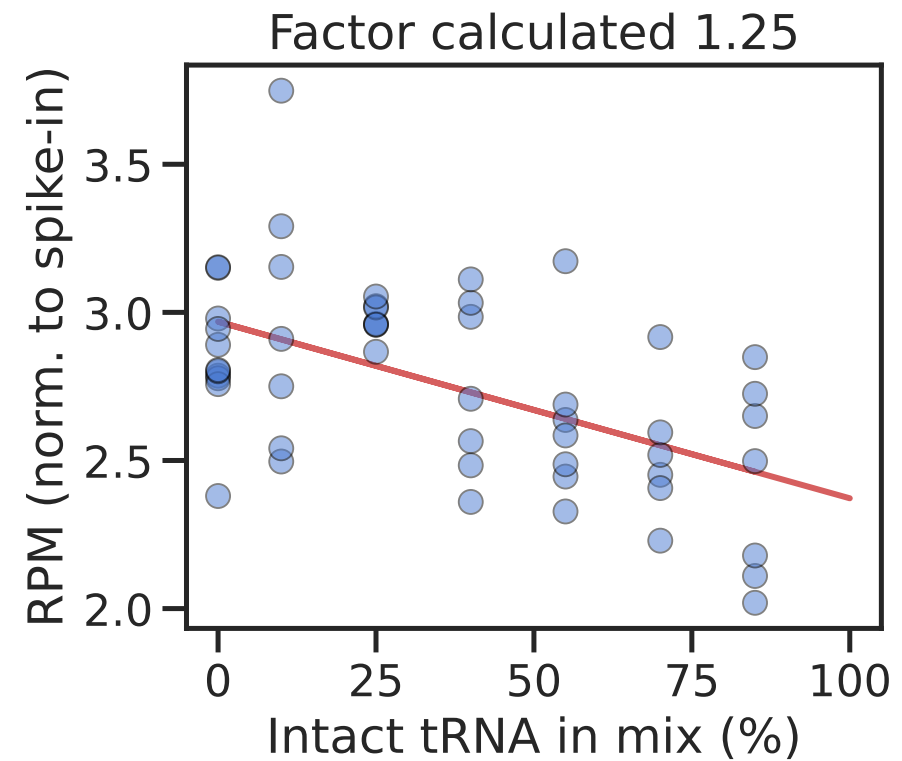
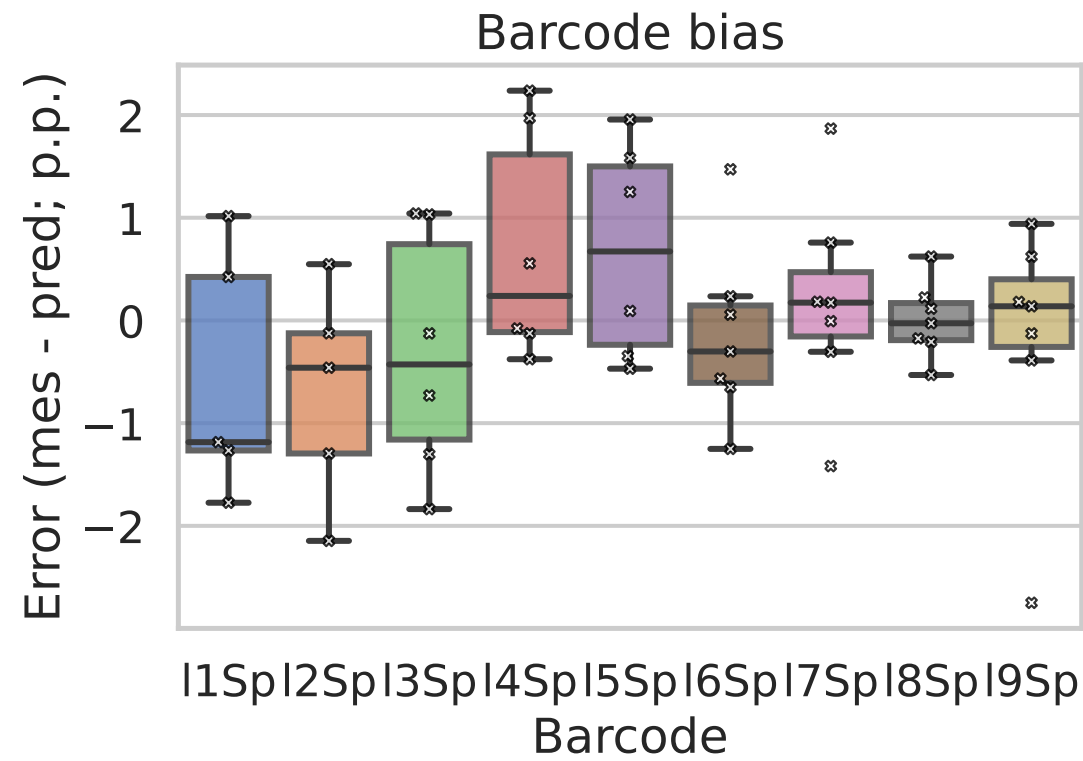
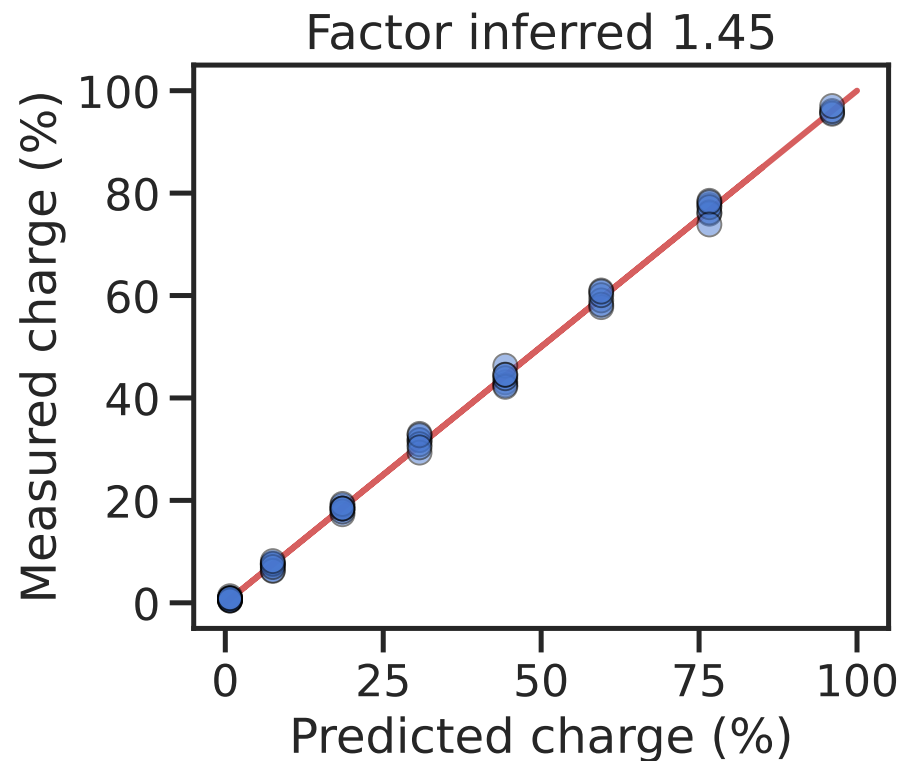
Leu-TAG-3-1



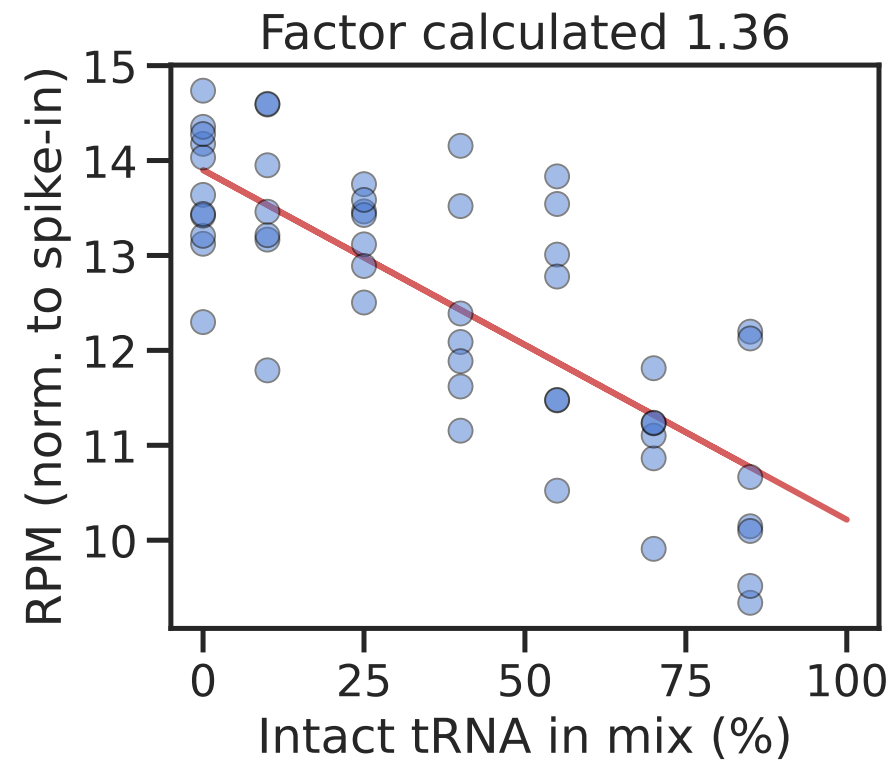
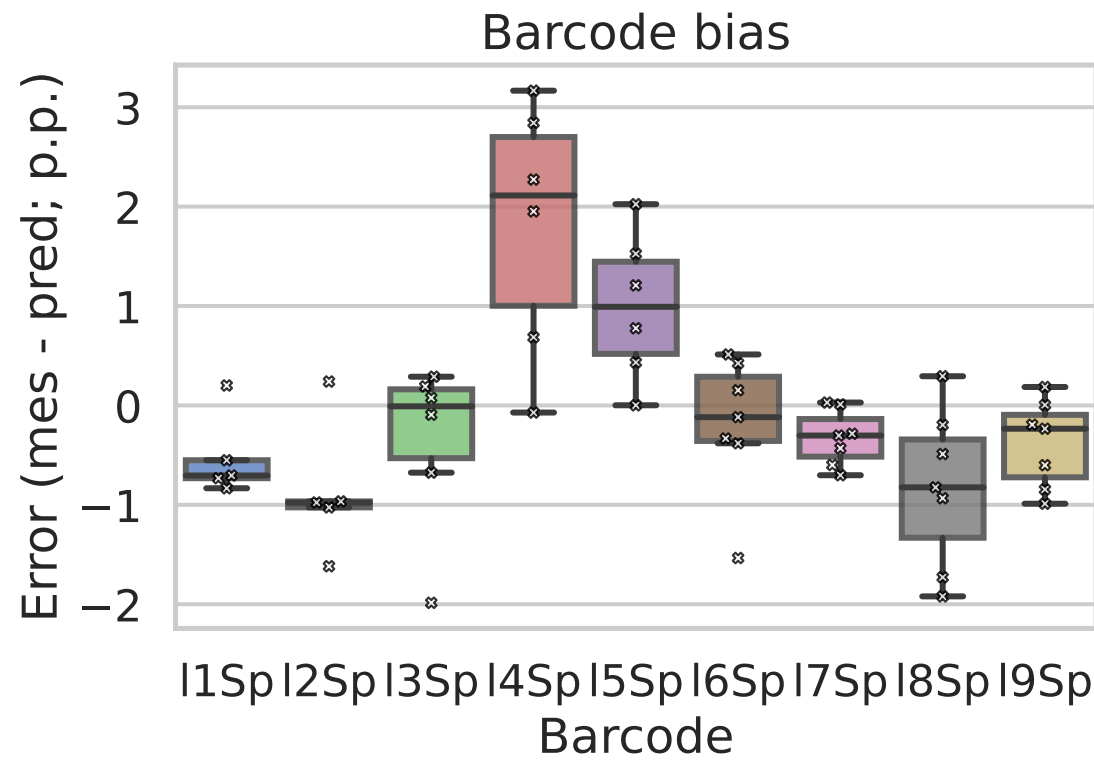
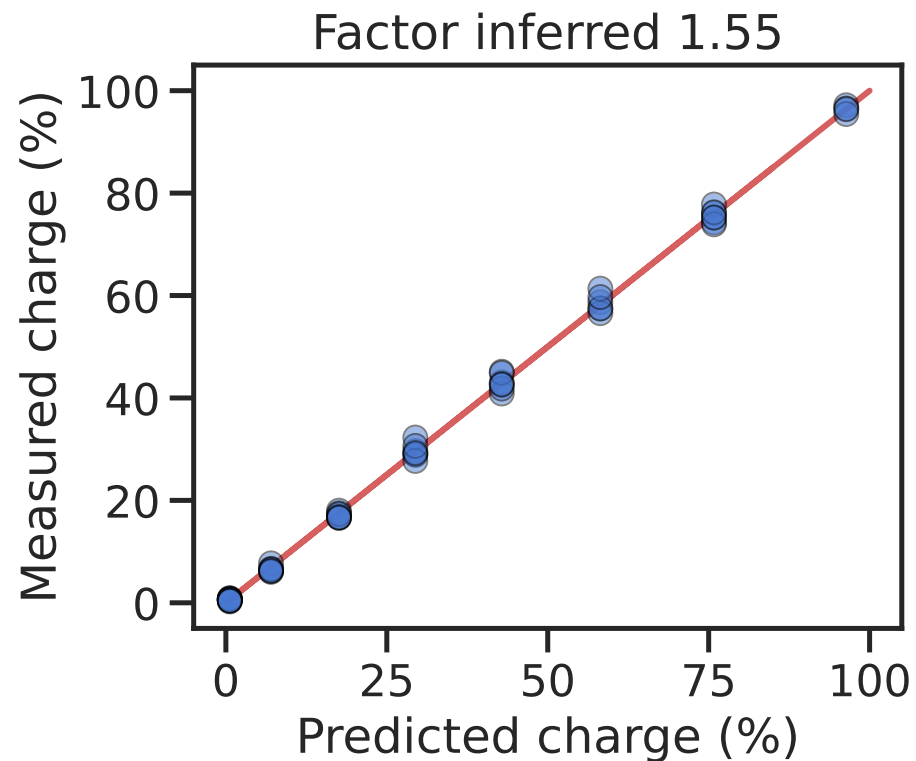
Pro-TGG-3-2



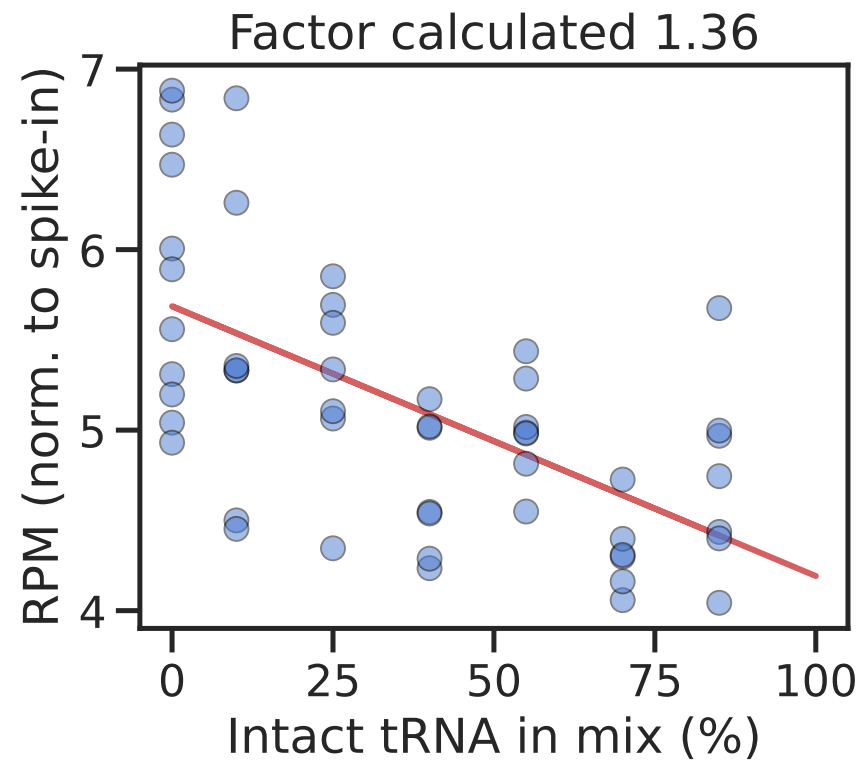
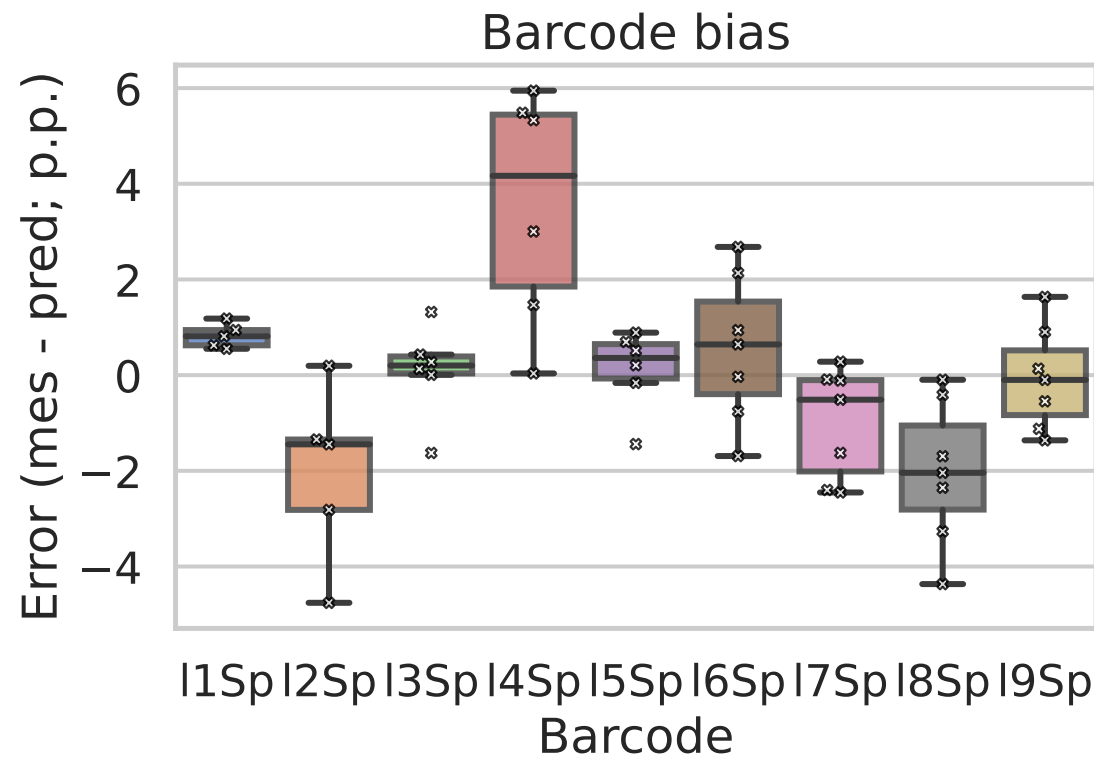
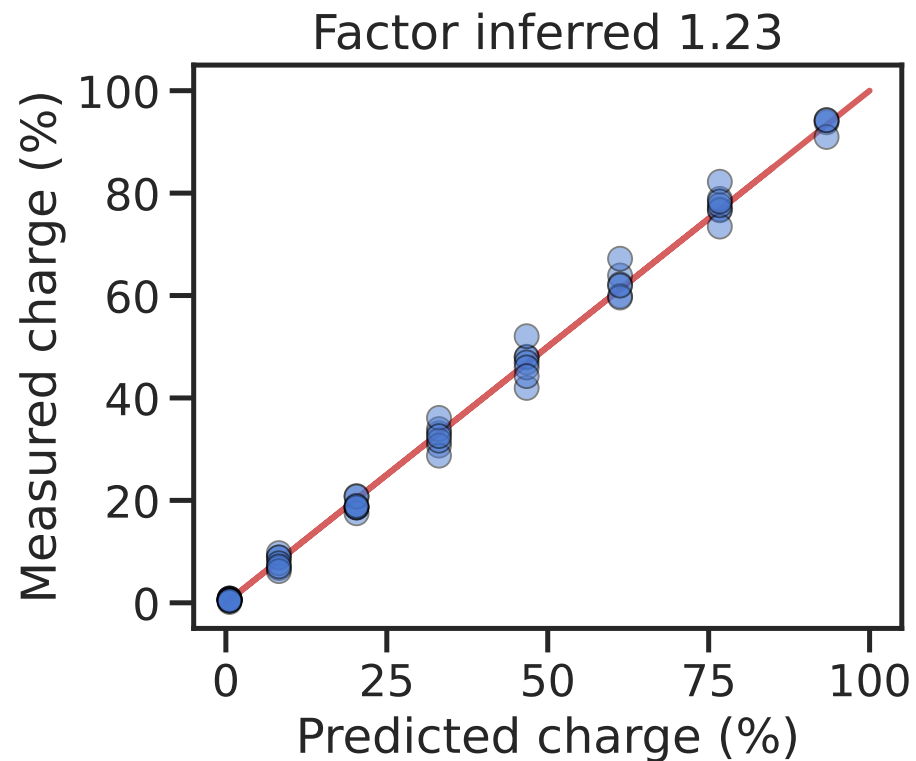
mito-Ala-TGC



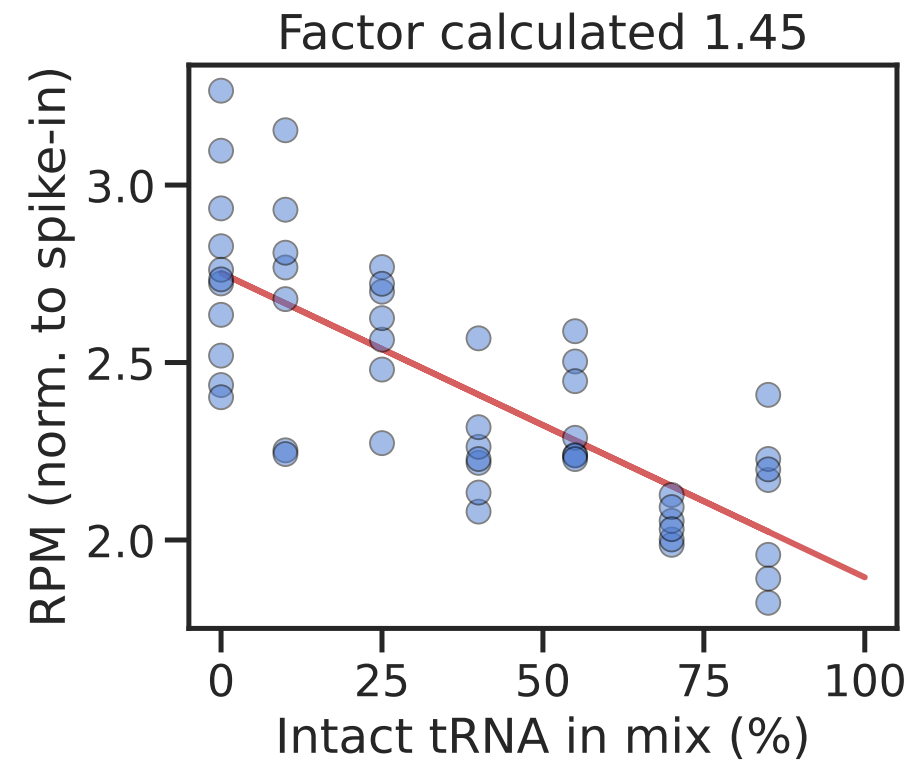
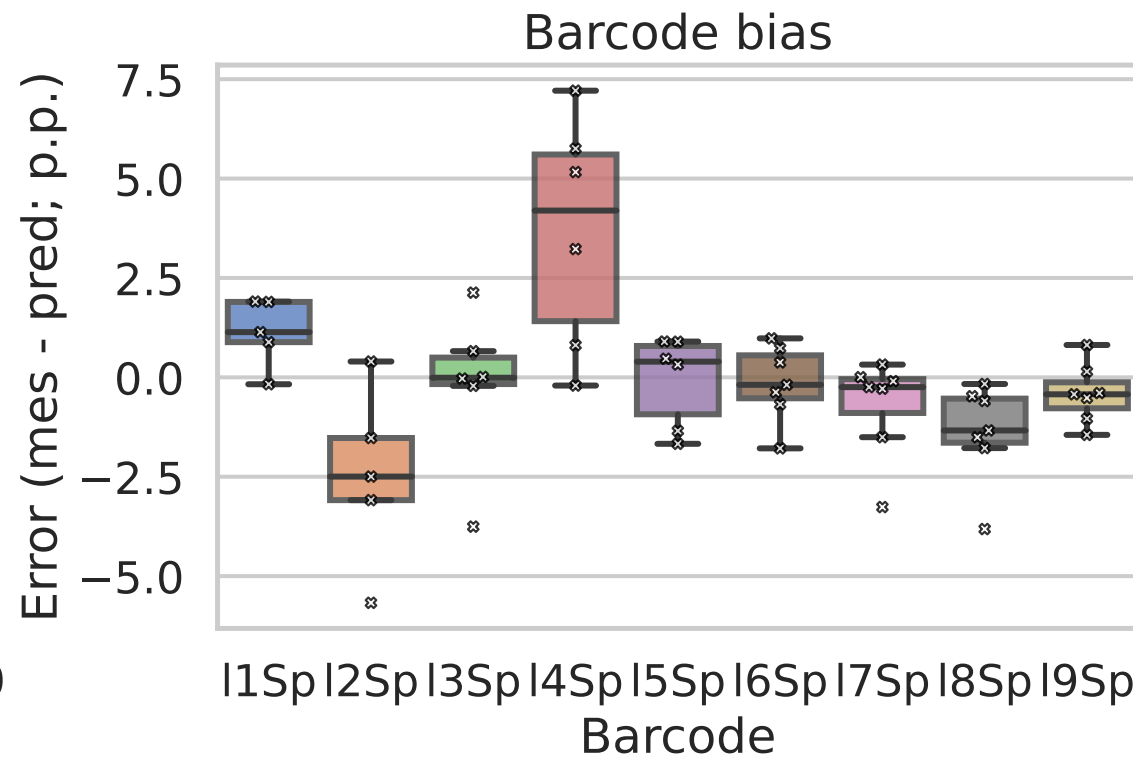
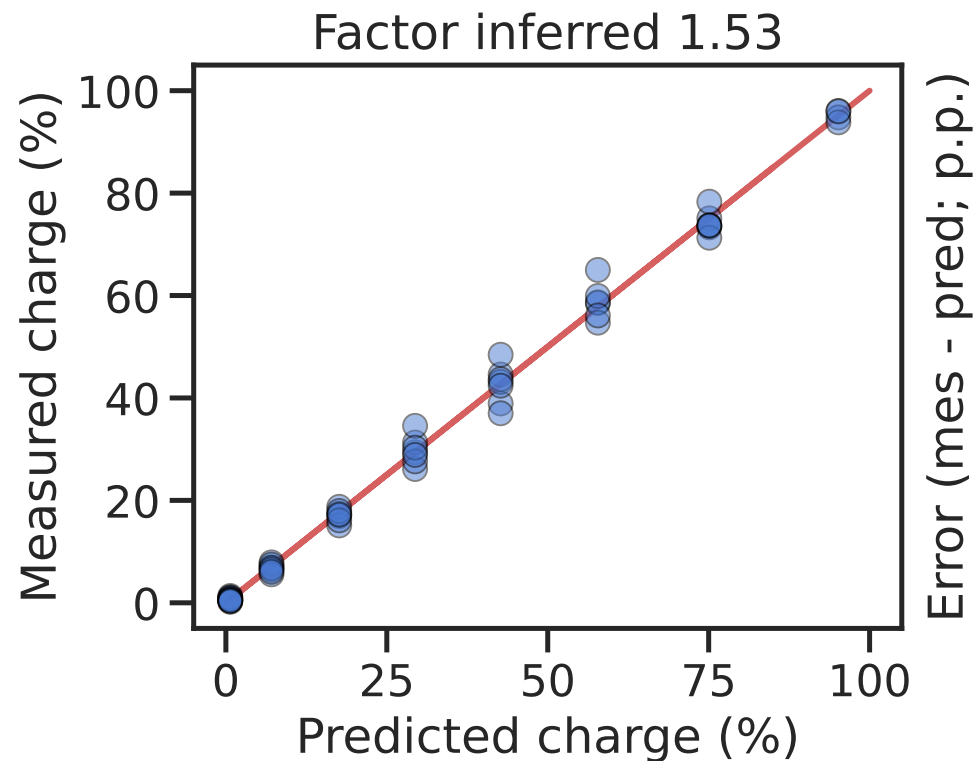
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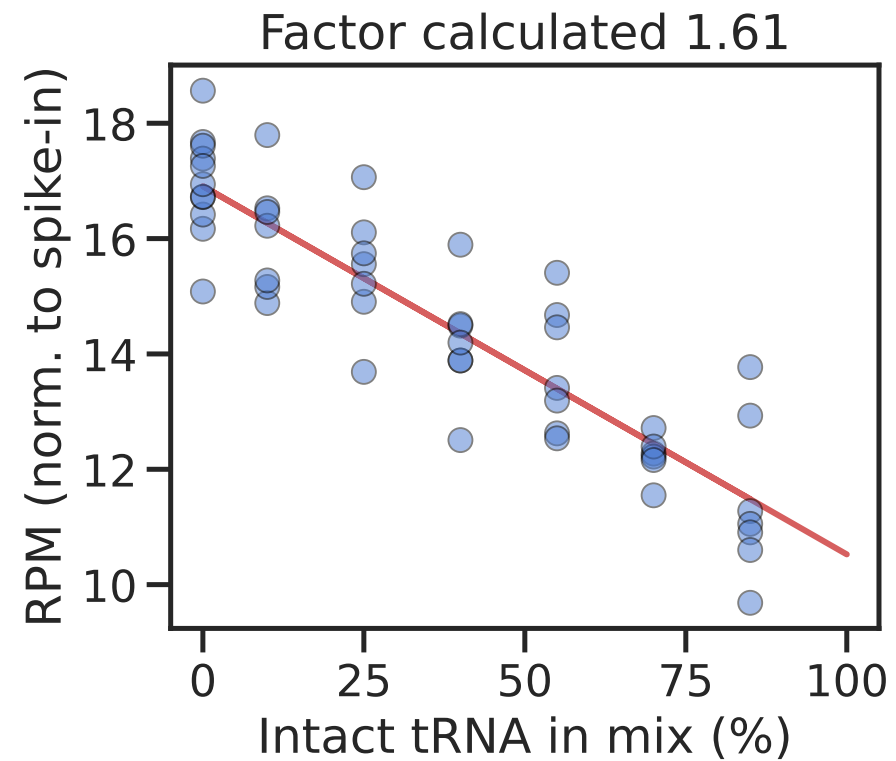
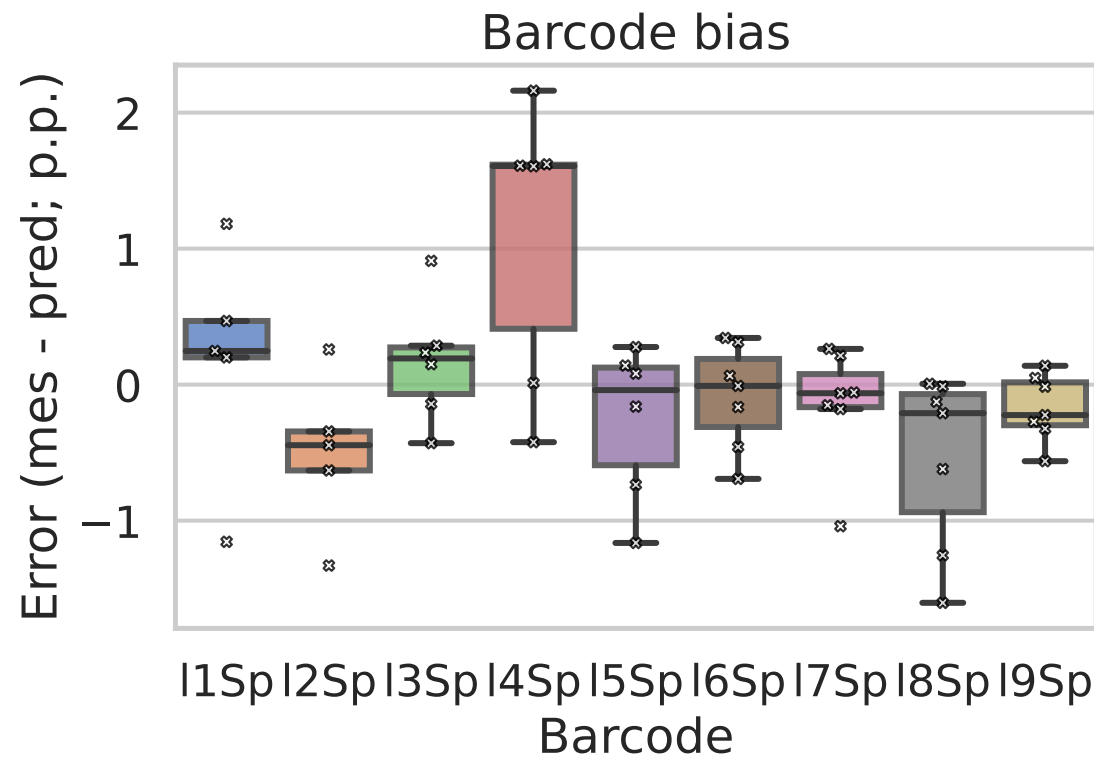
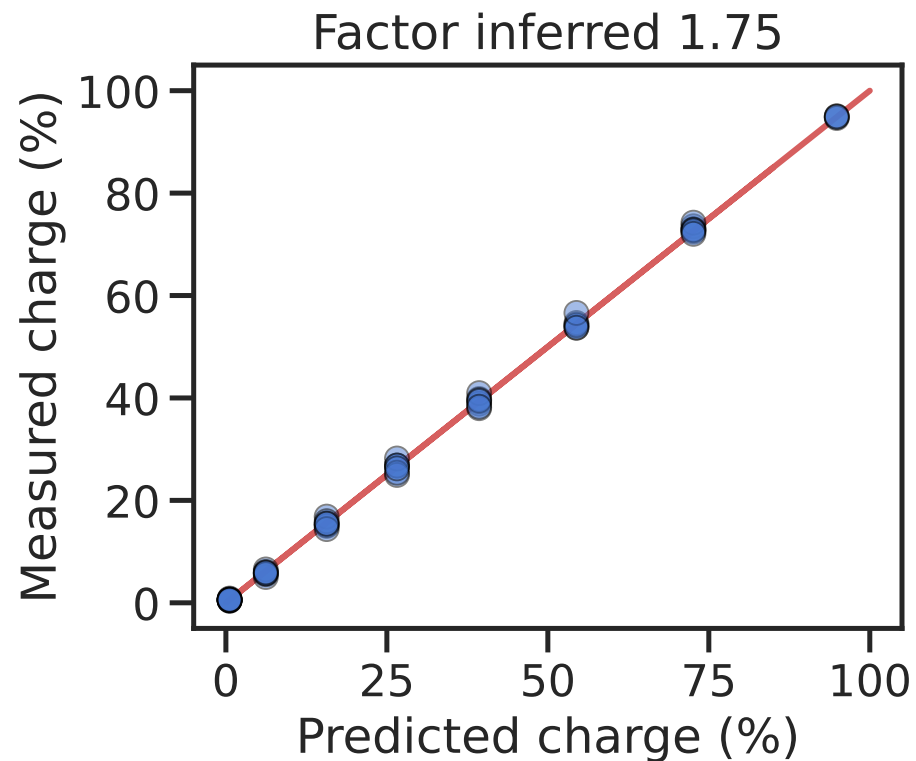
Thr-TGT-4-1



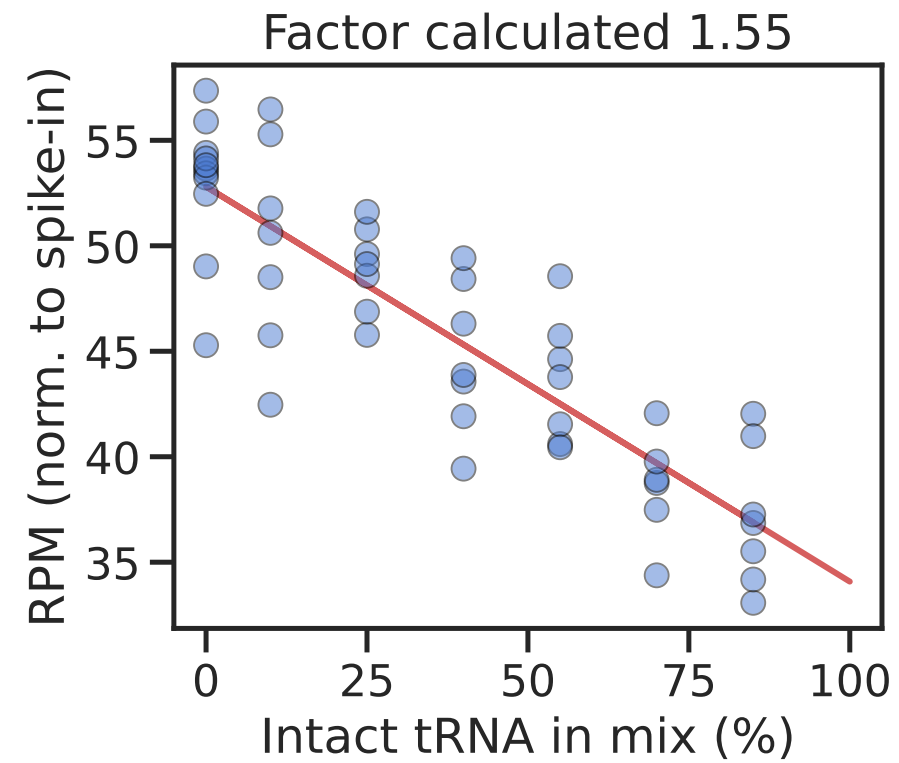
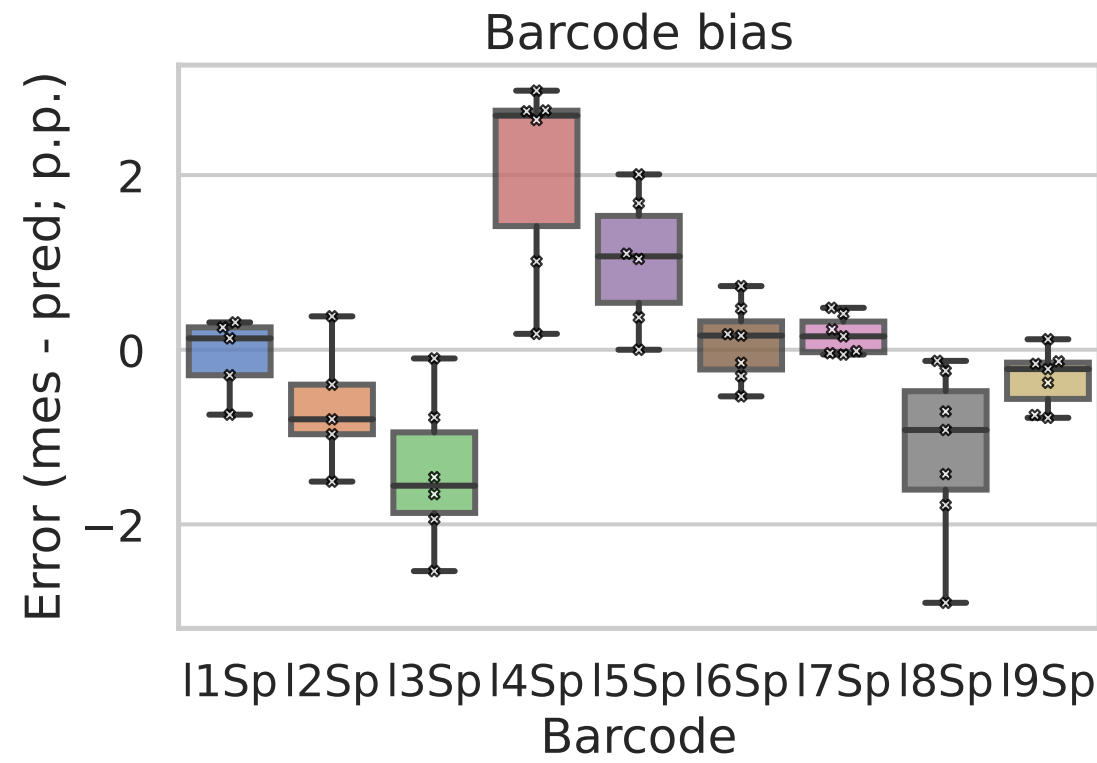
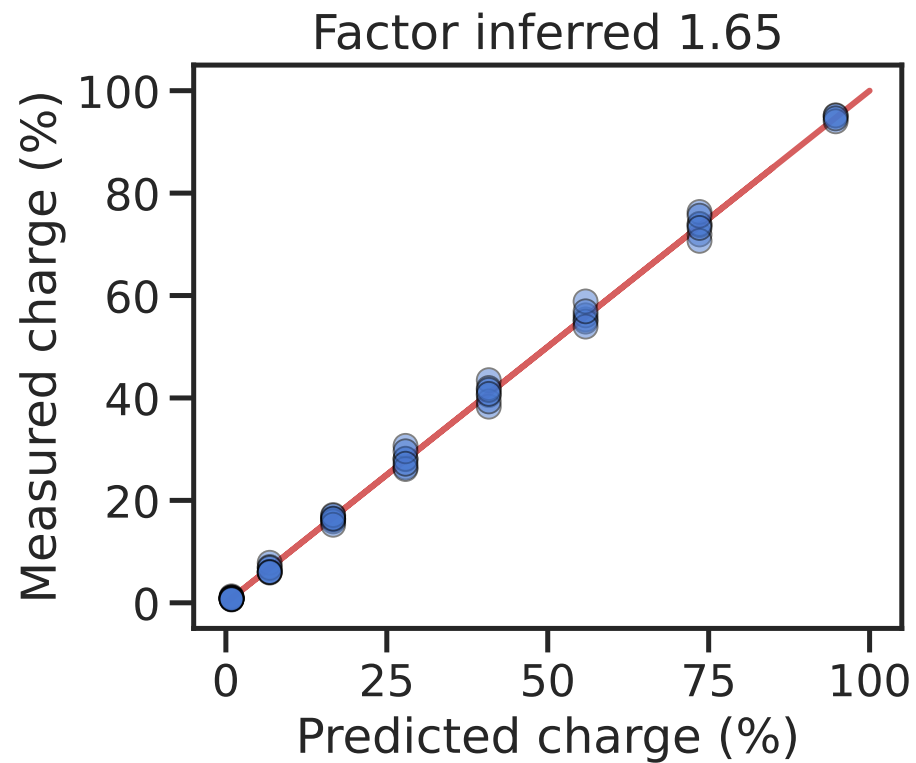
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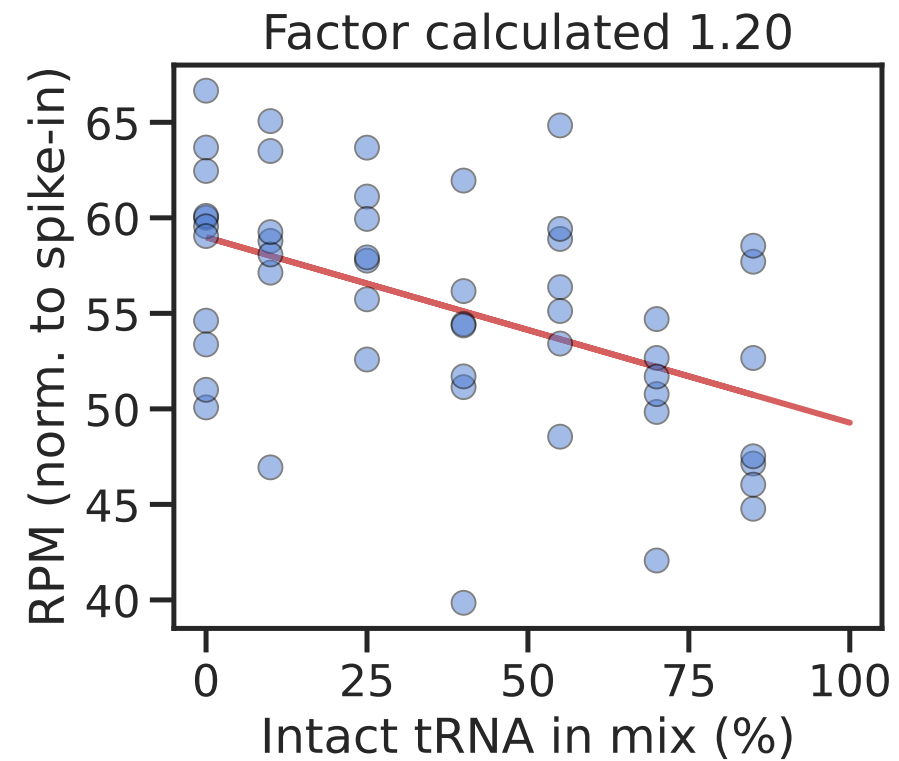
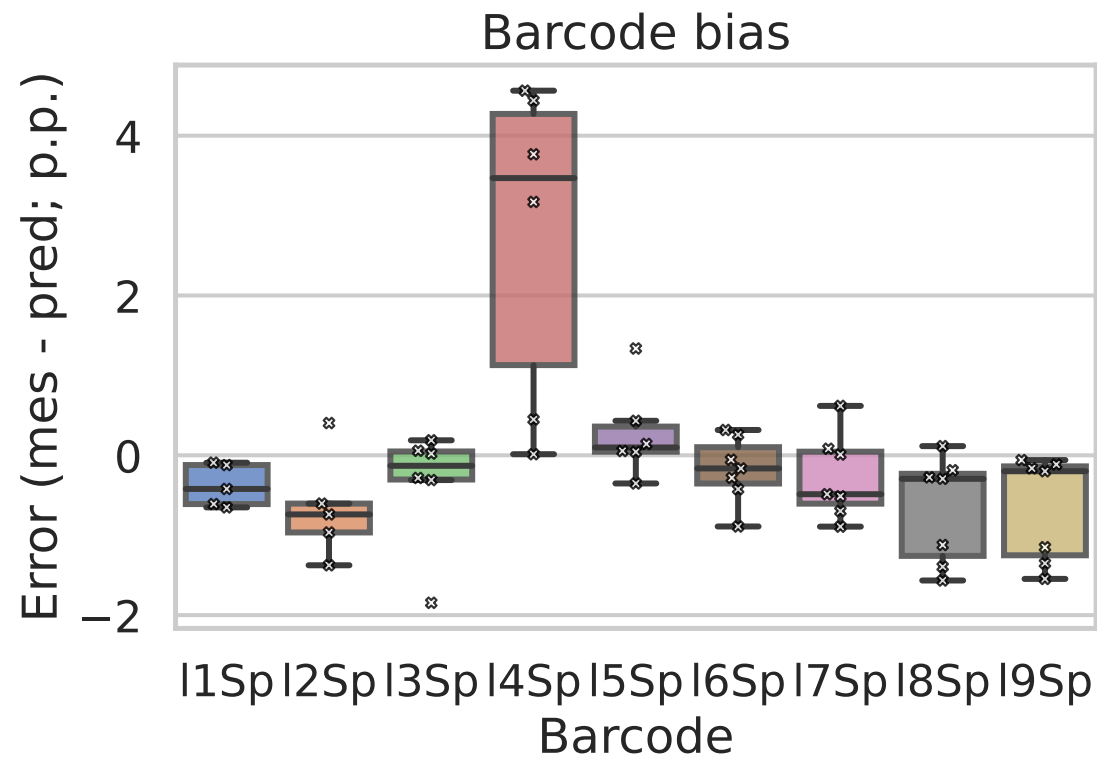
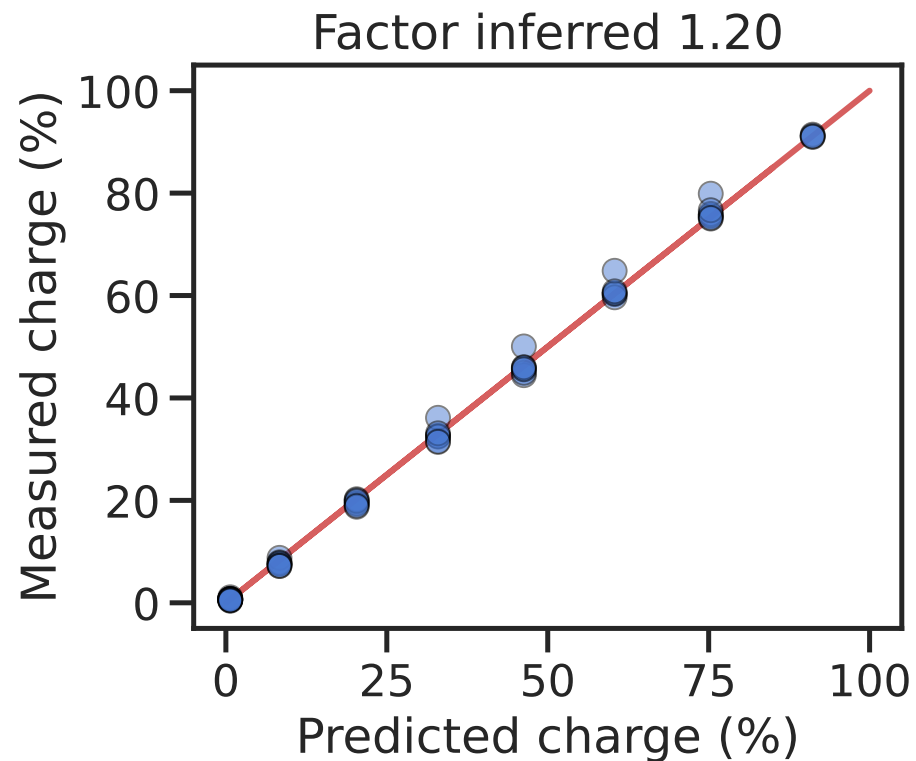
Ala-AGC-11-1



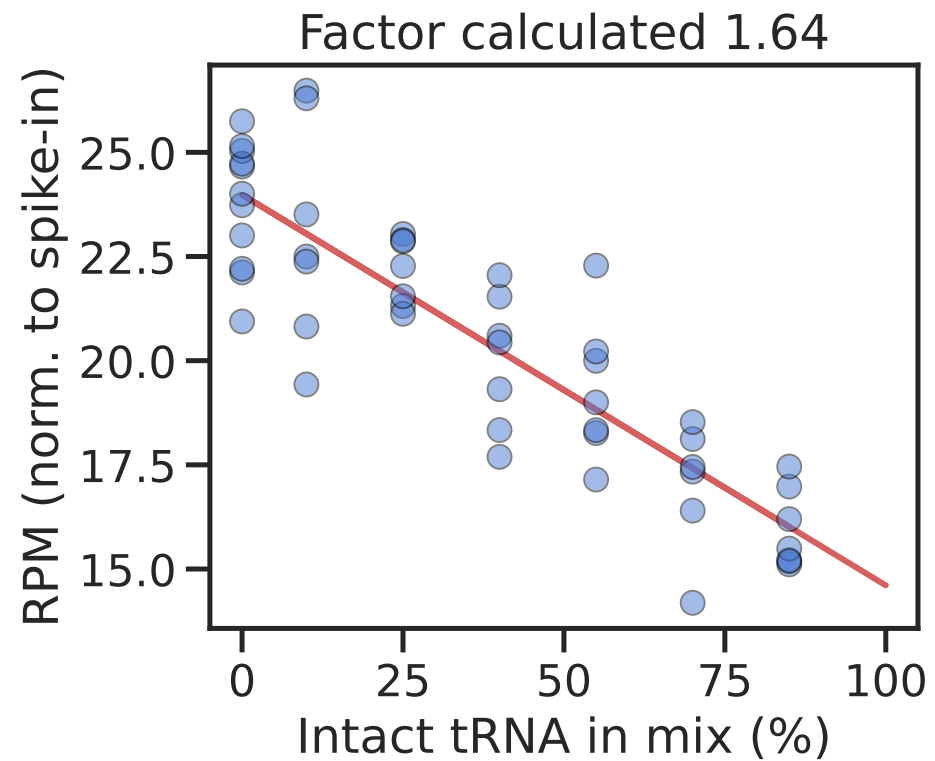
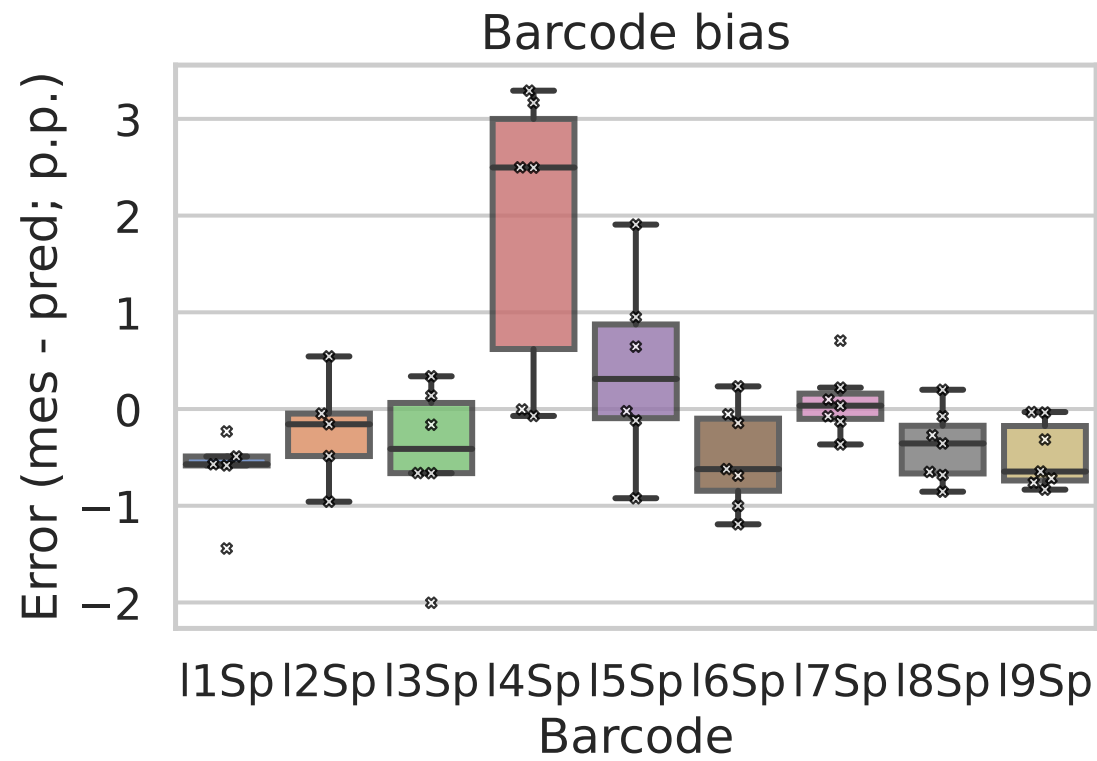
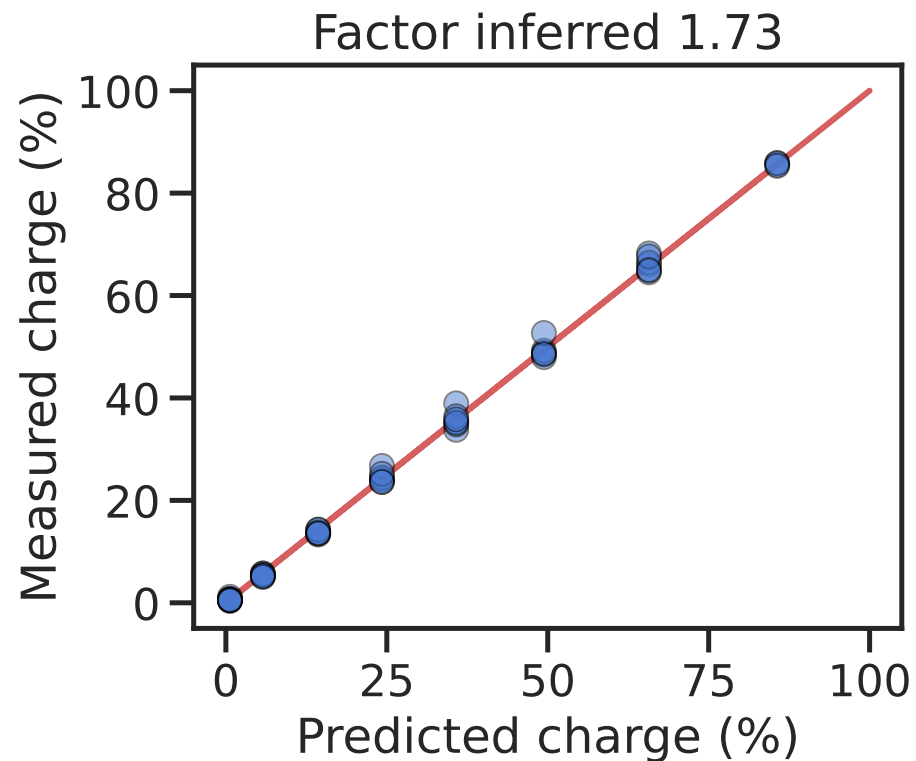
Gly-GCC-2-1



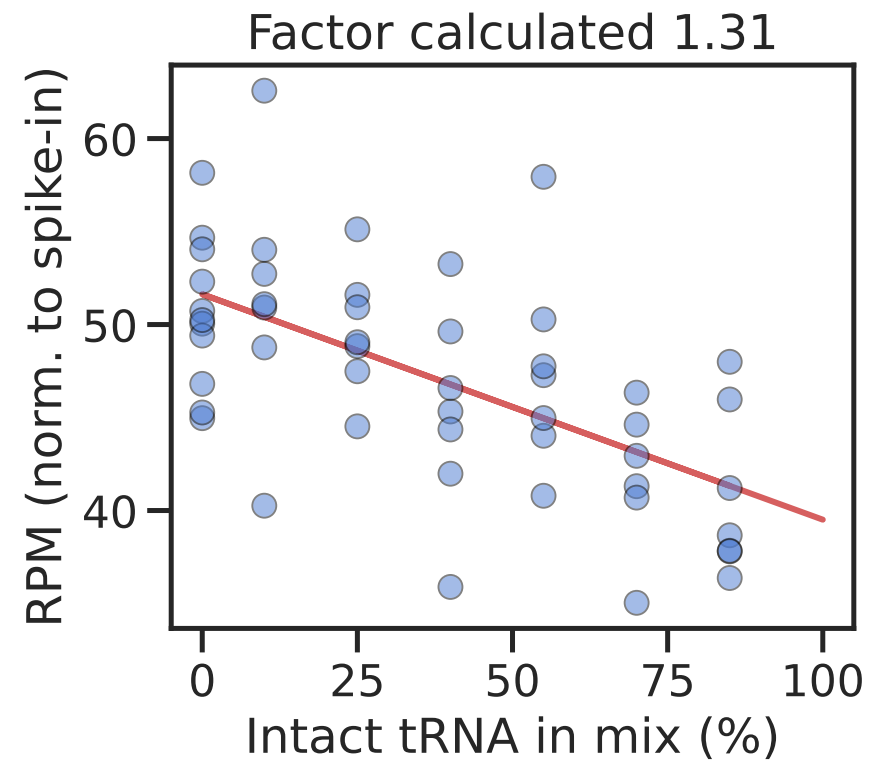
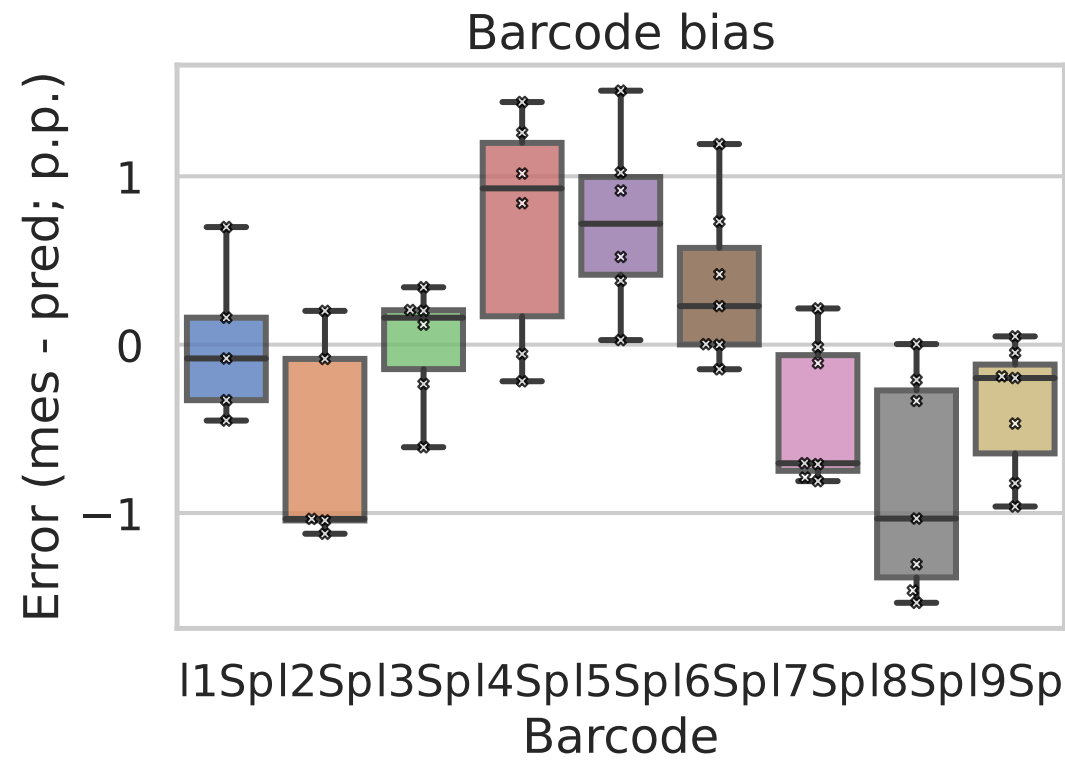
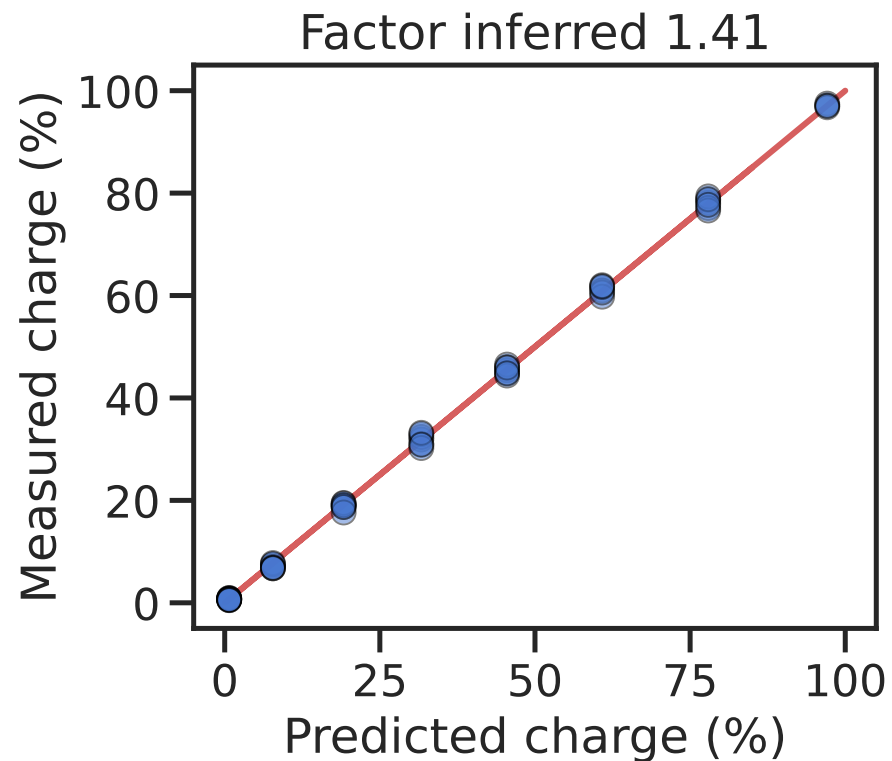
Glu-CTC-1-1



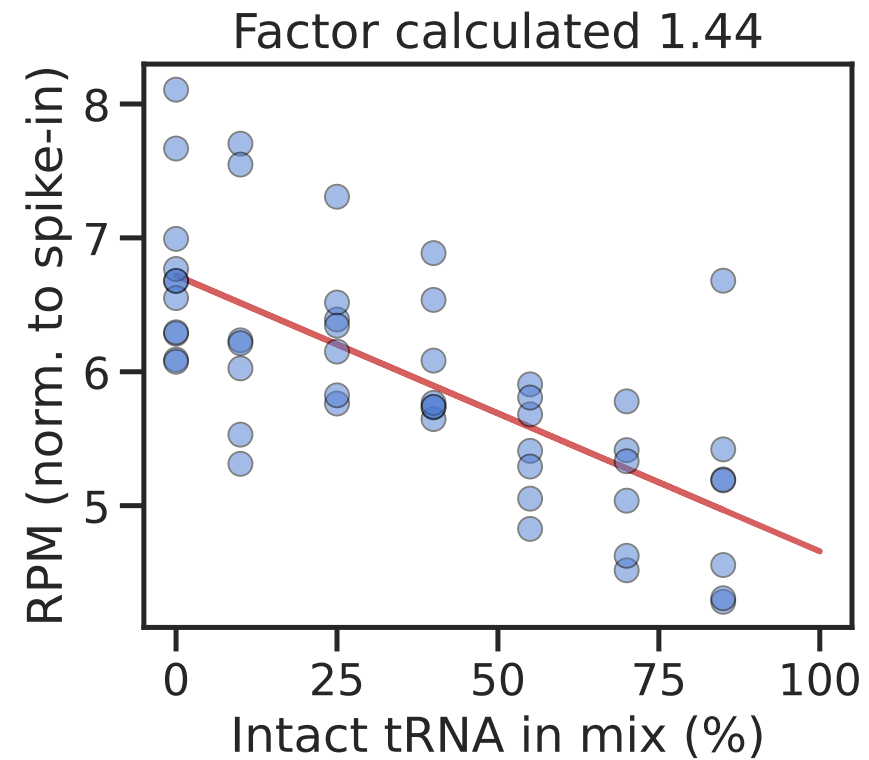
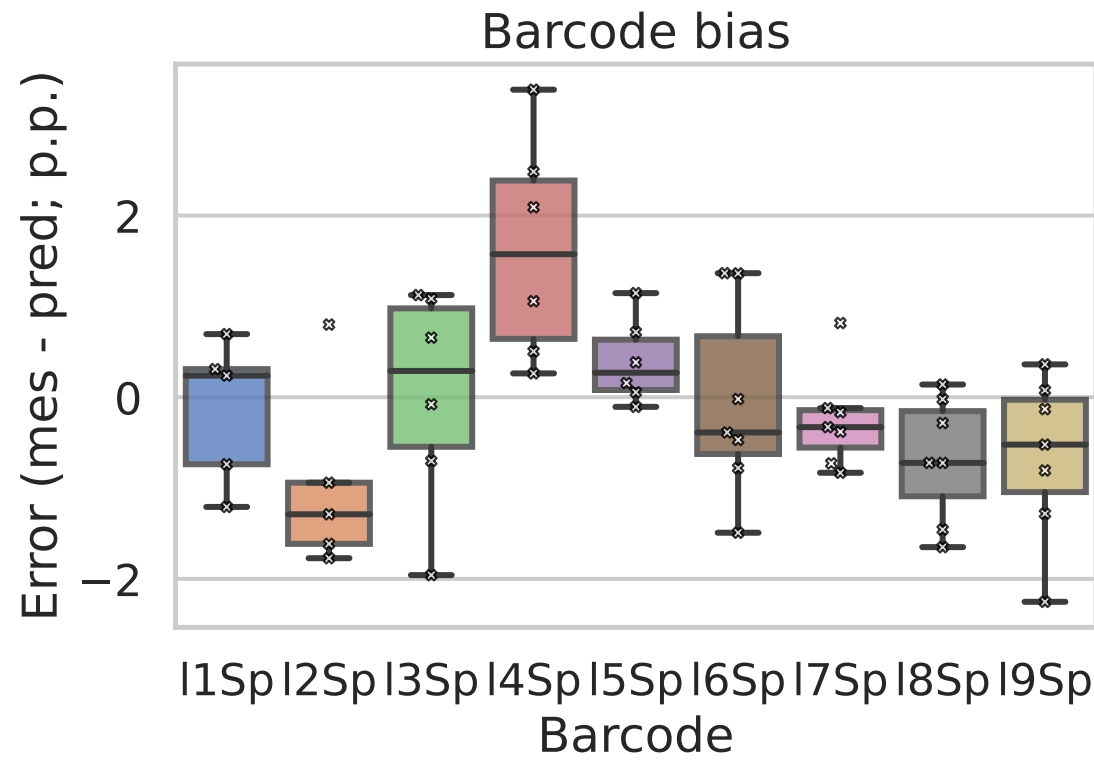
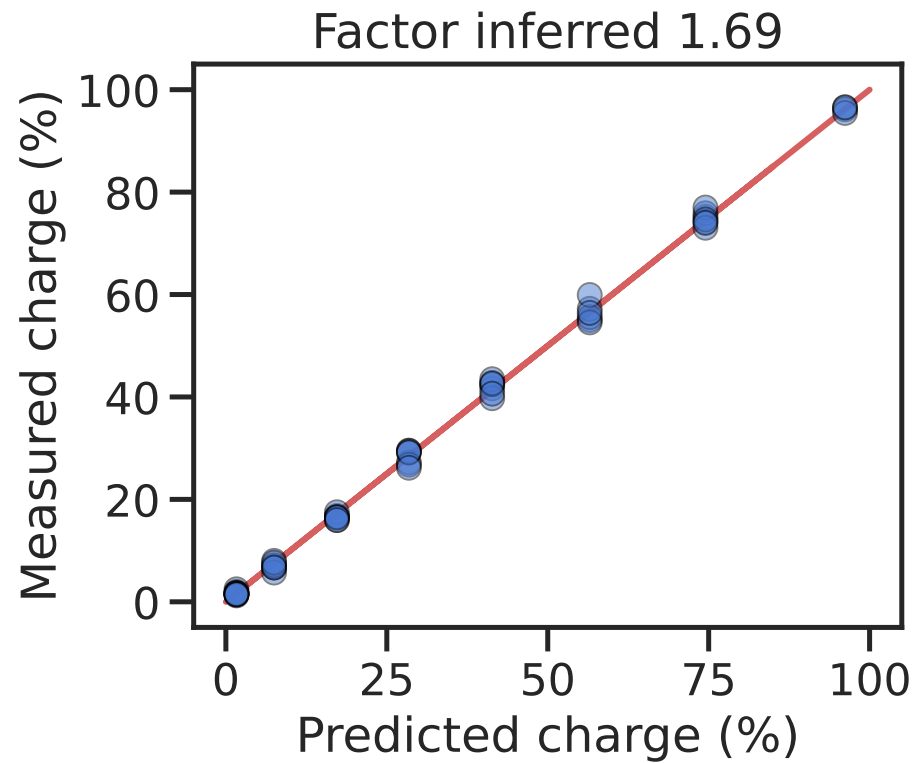
Glu-TTC-1-2



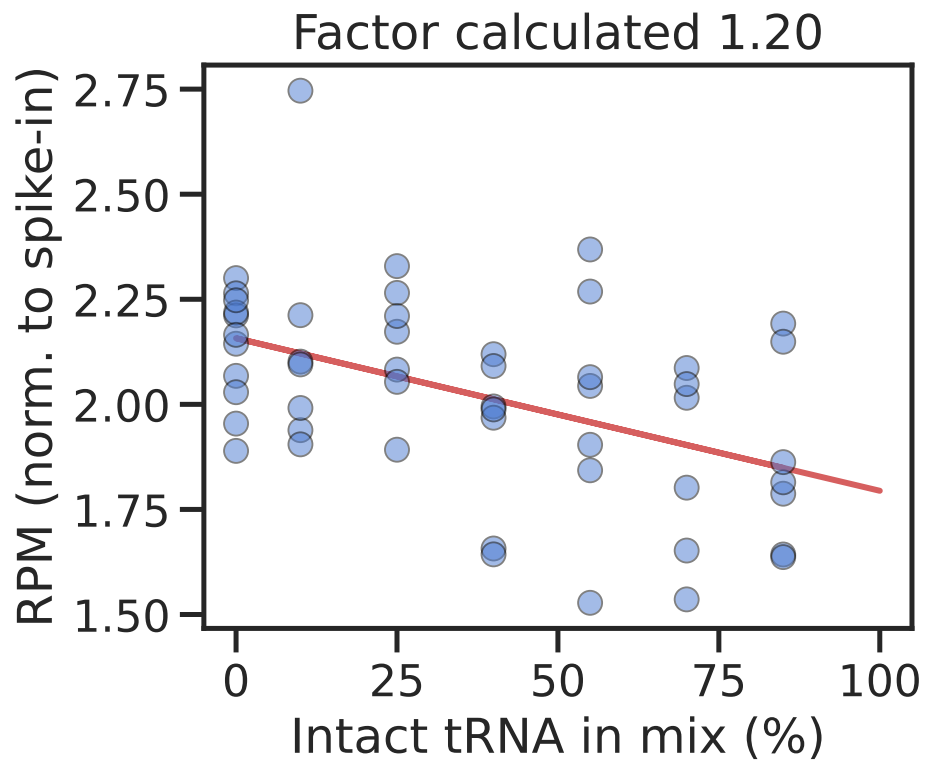
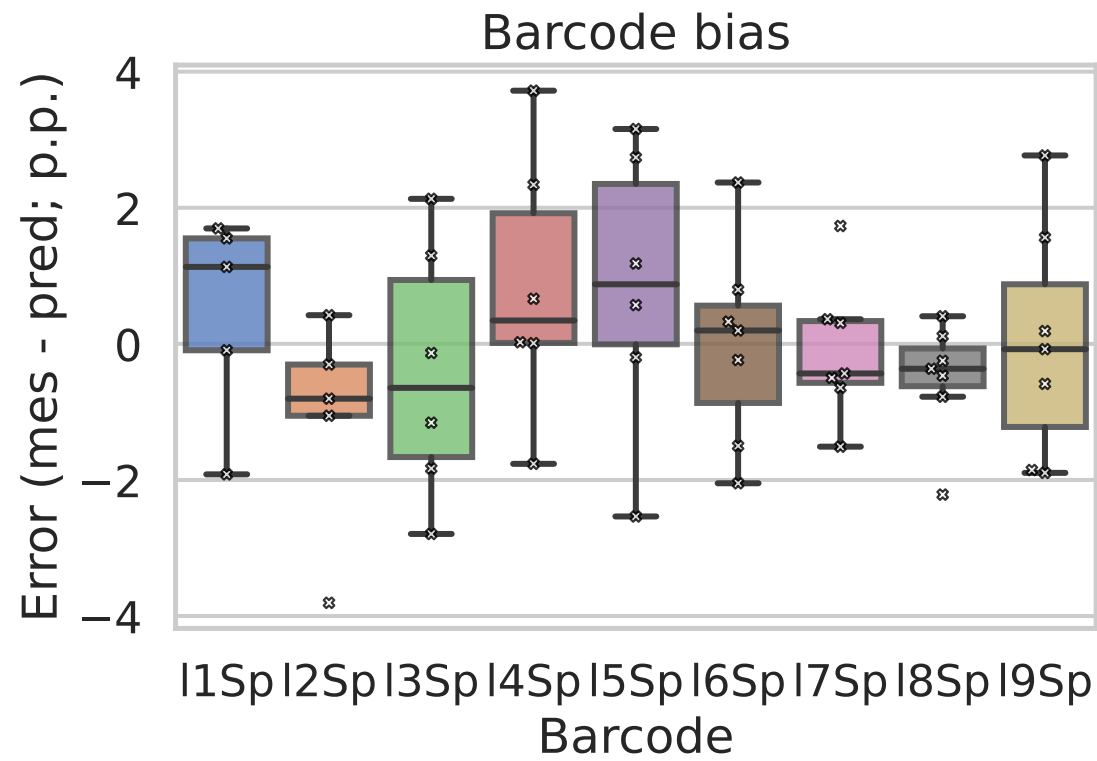
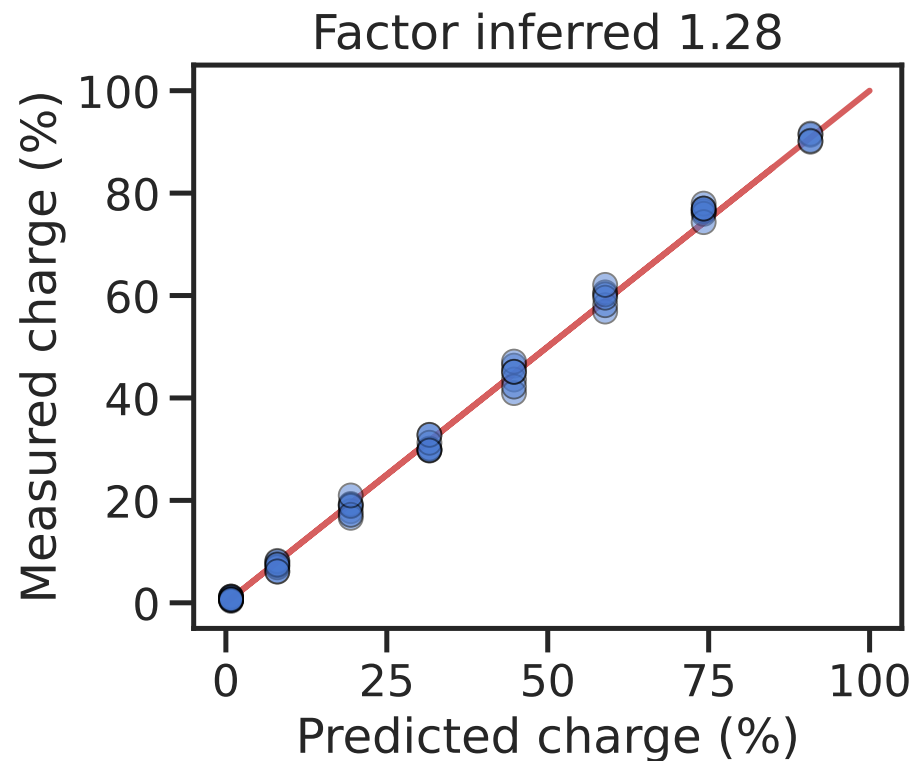
Lys-CTT-2-1



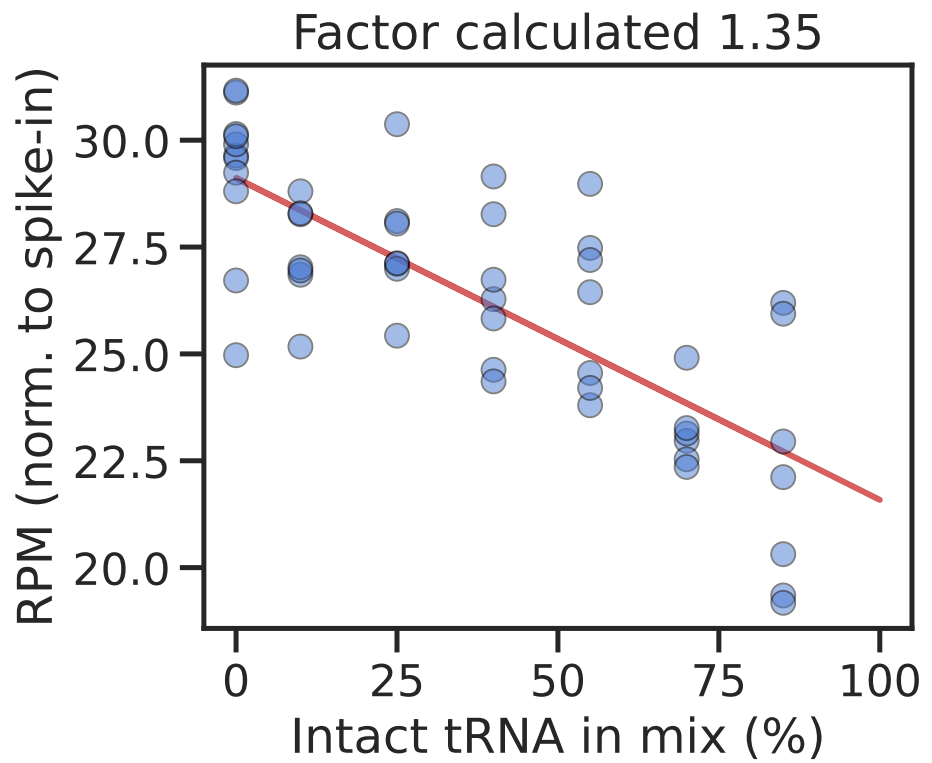
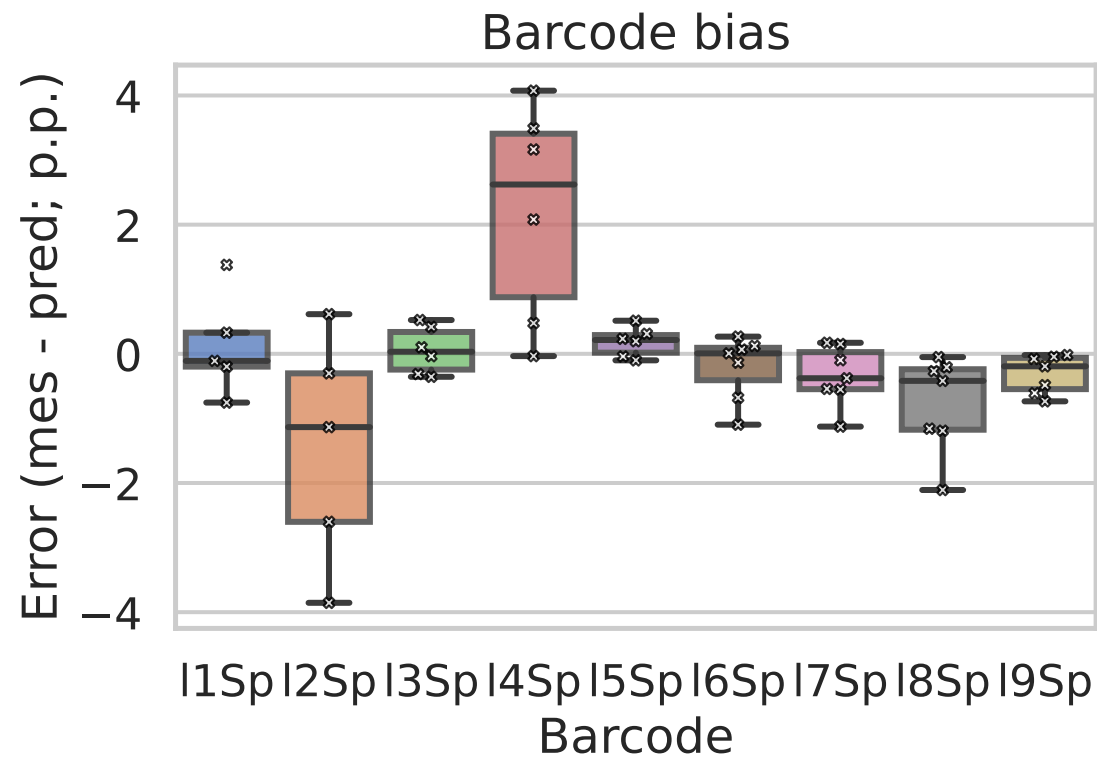
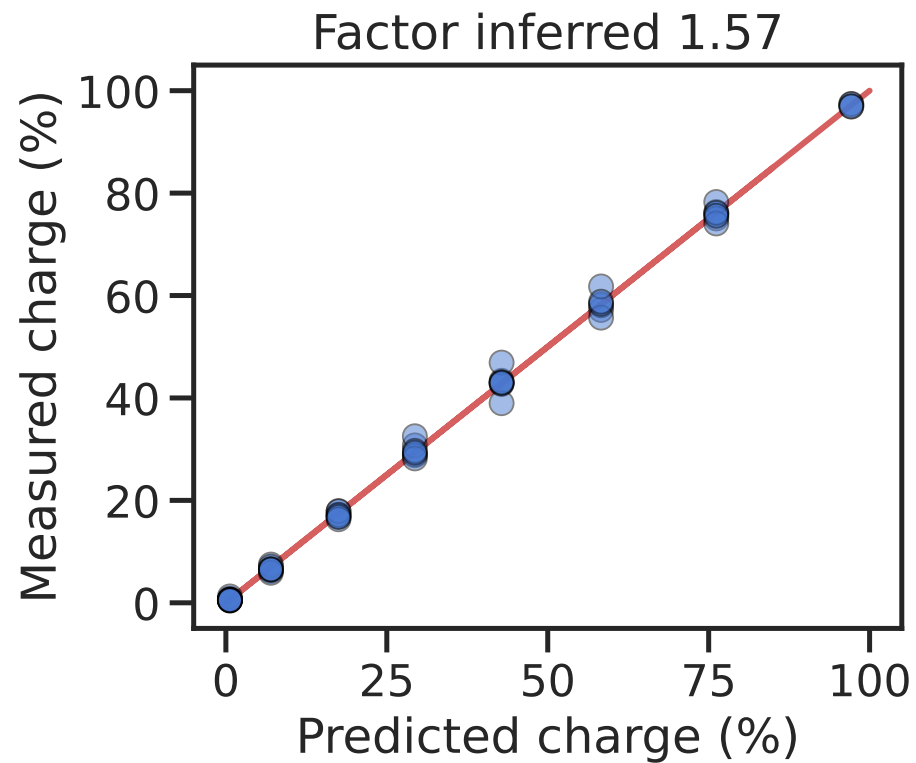
mito-Trp-TCA



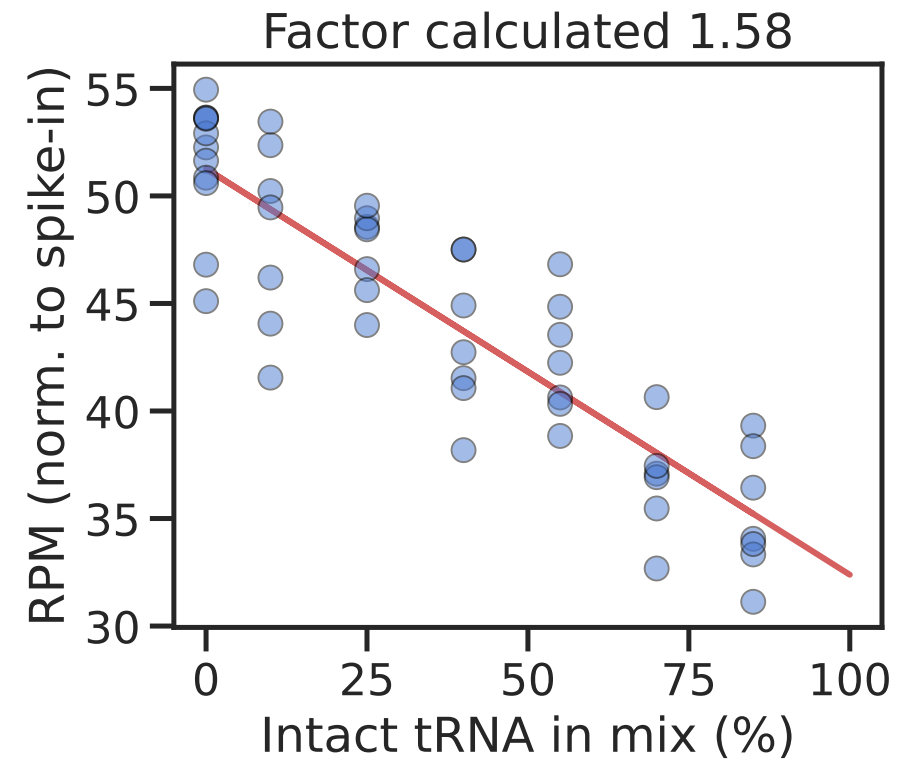
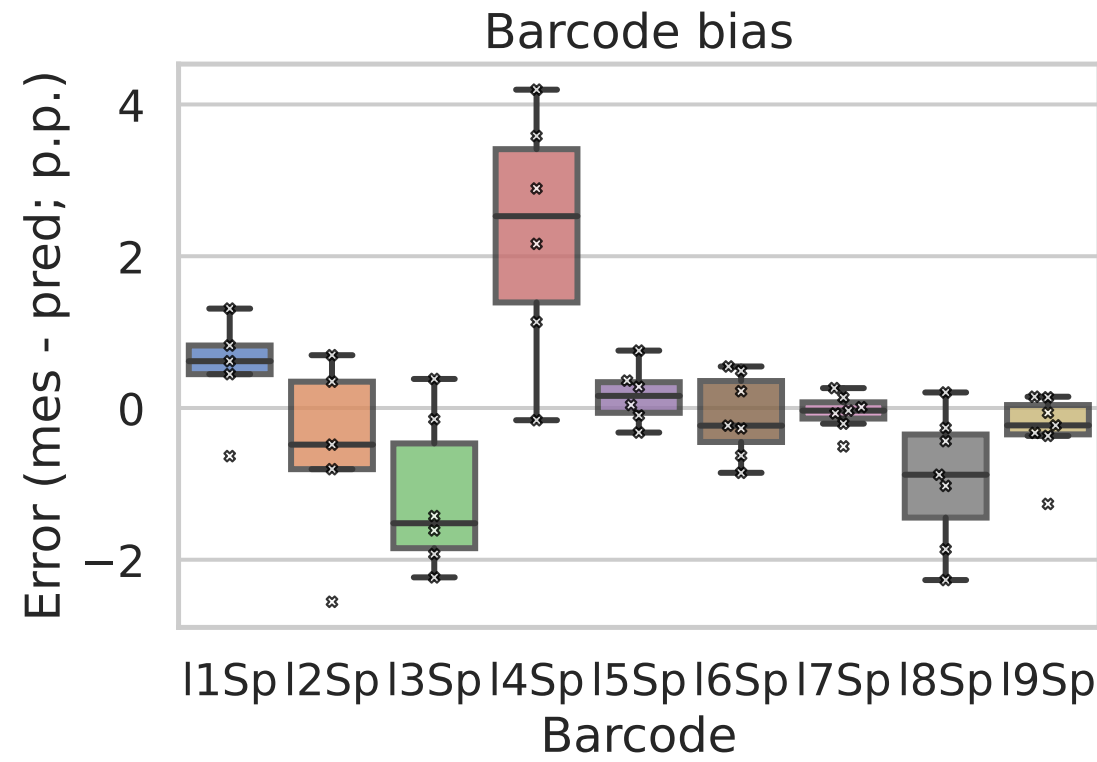
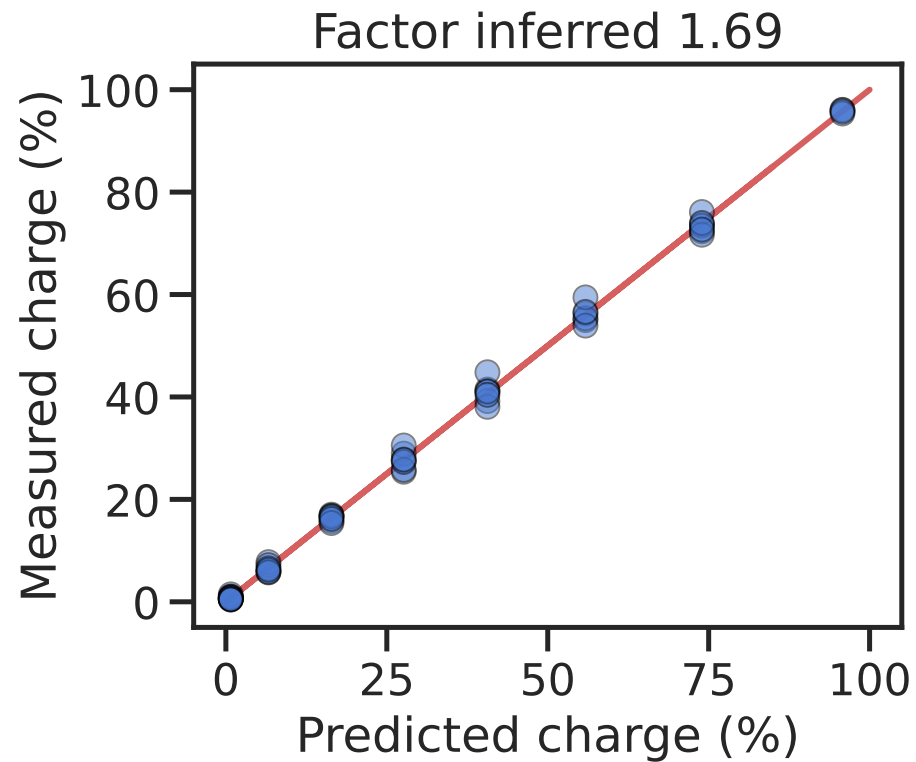
Ser-TGA-4-1



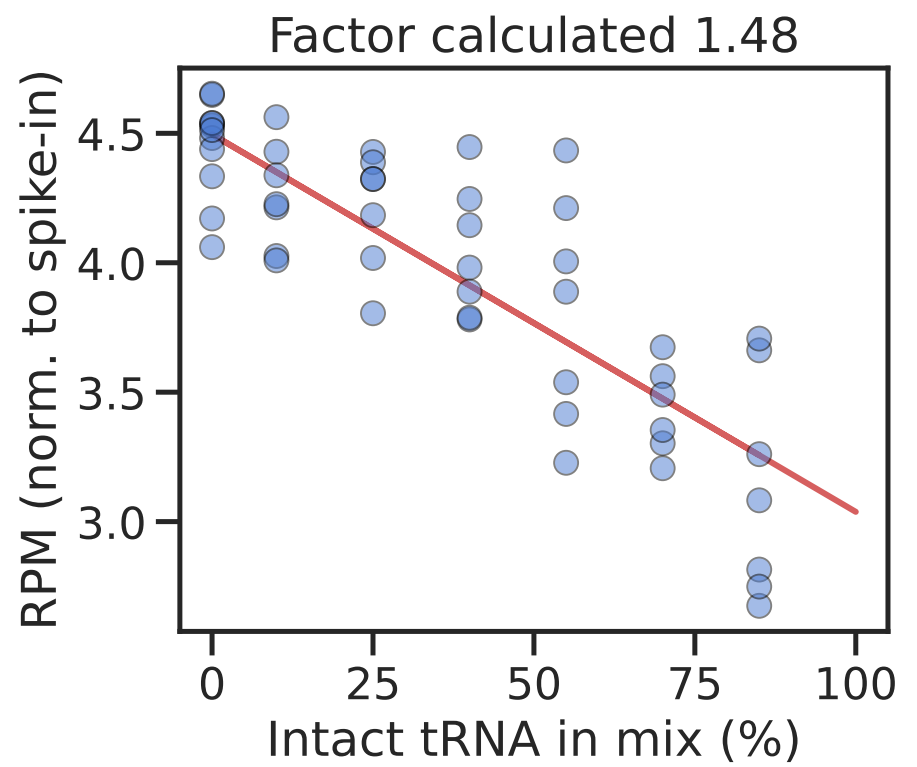
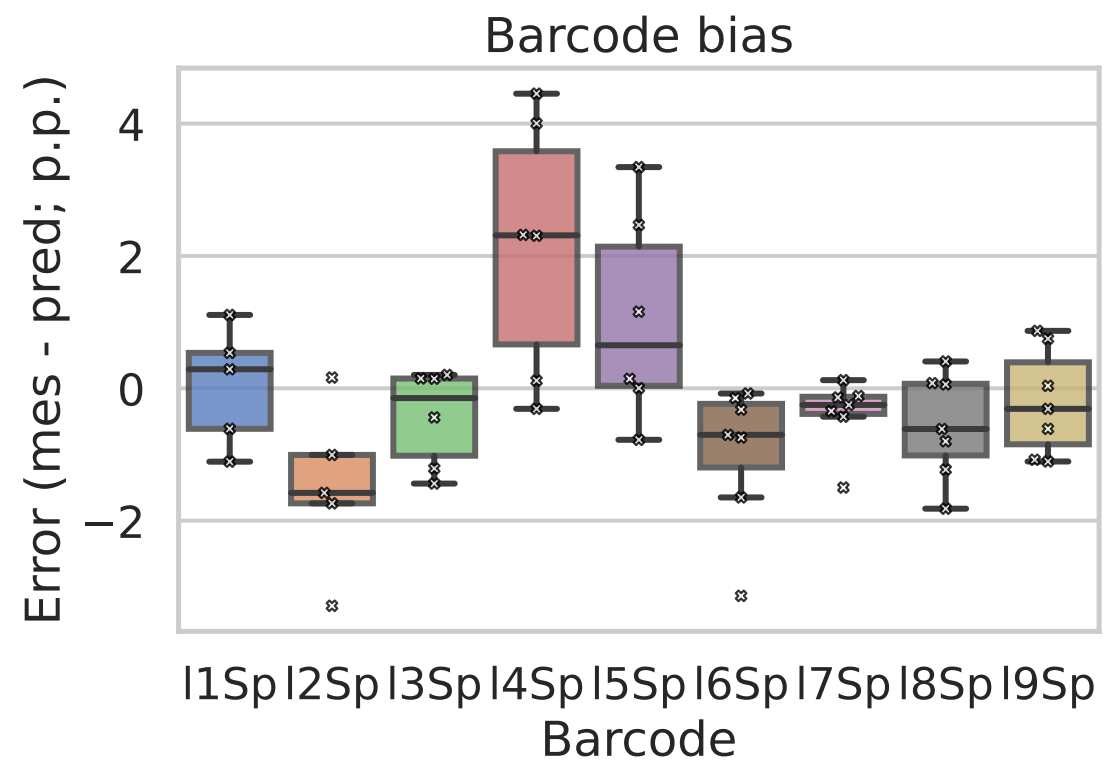
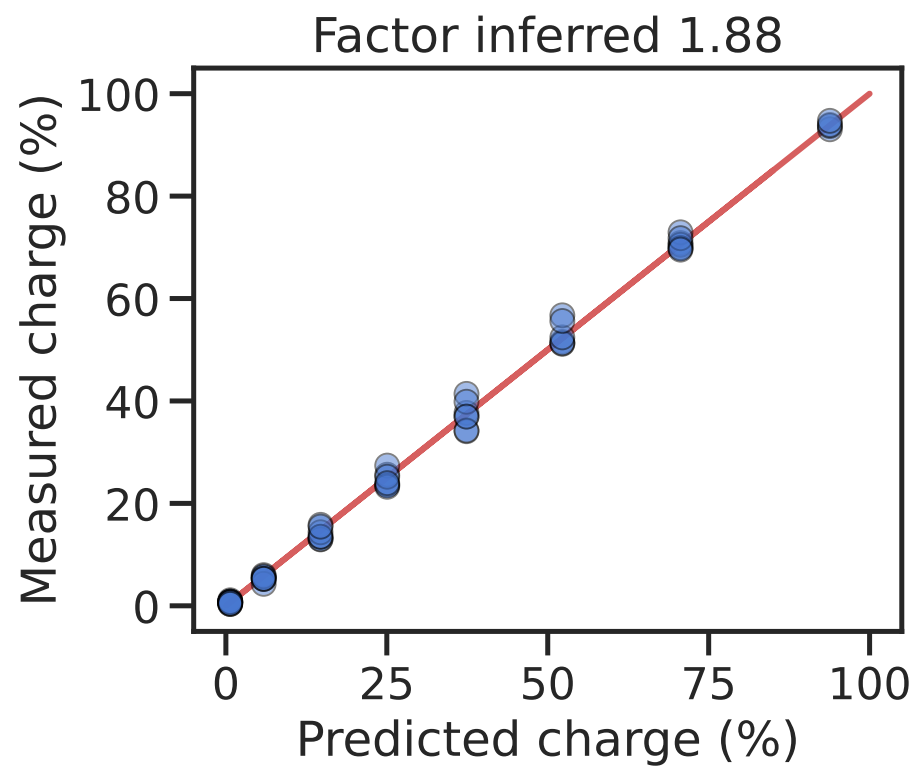
Ala-CGC-2-1



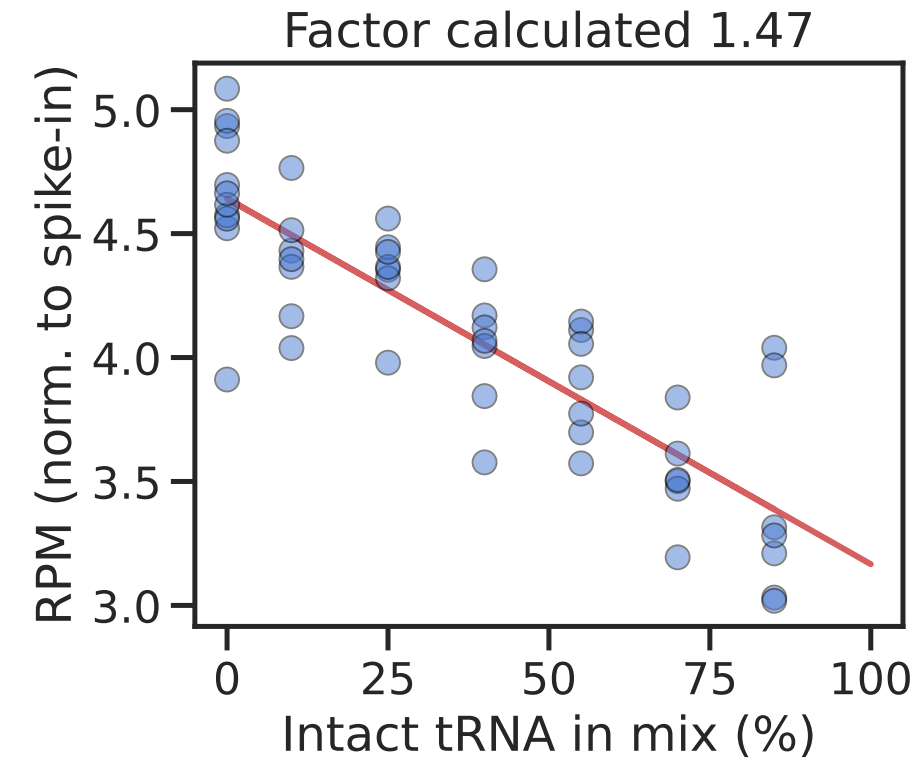
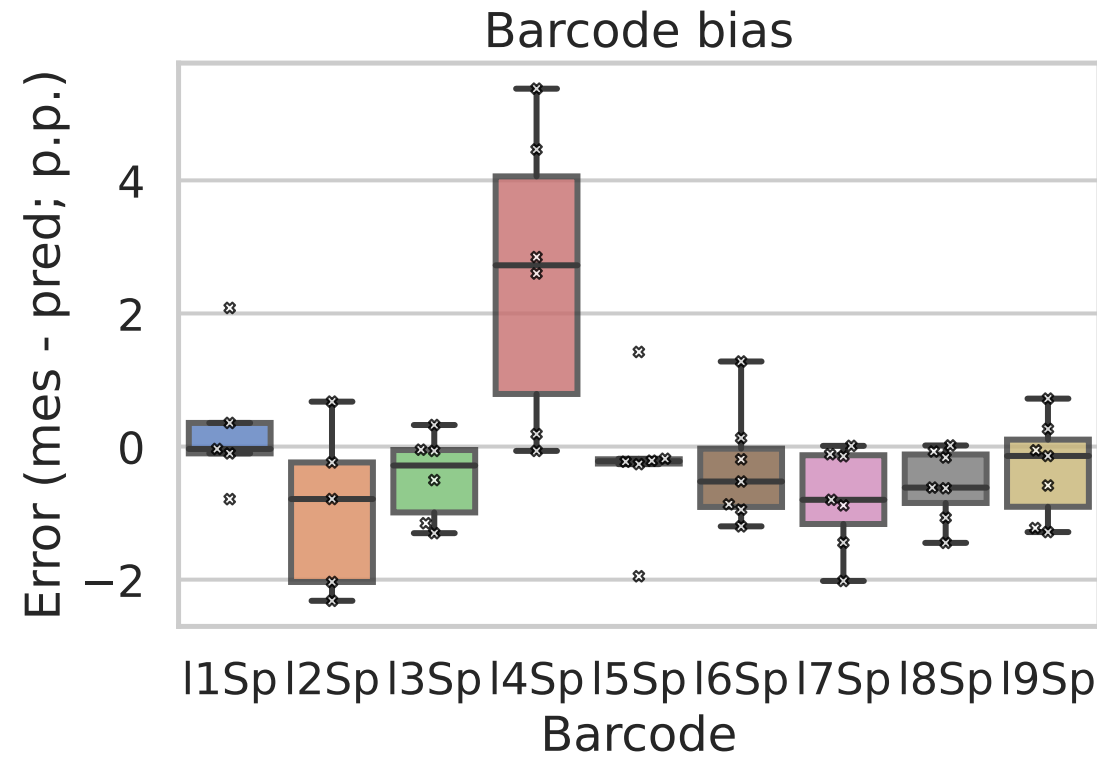
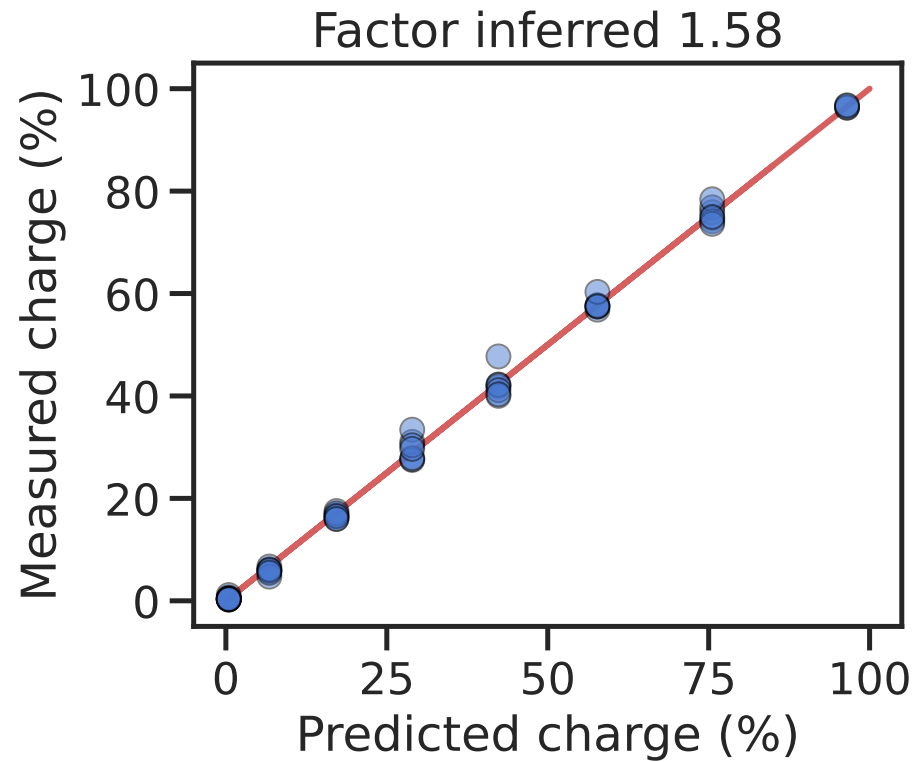
Gly-GCC-1-1



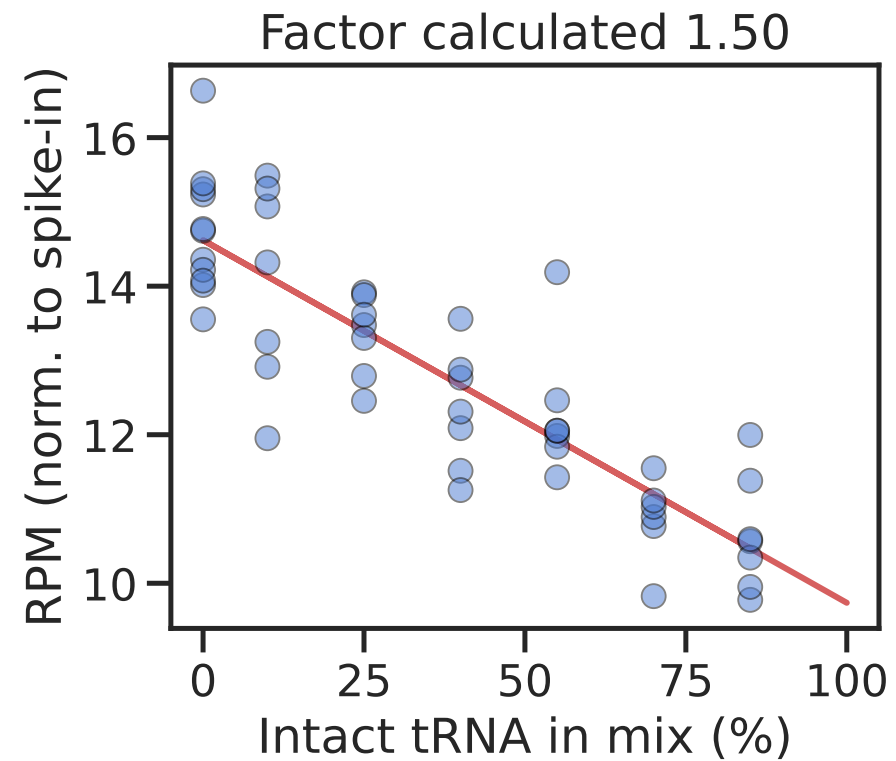
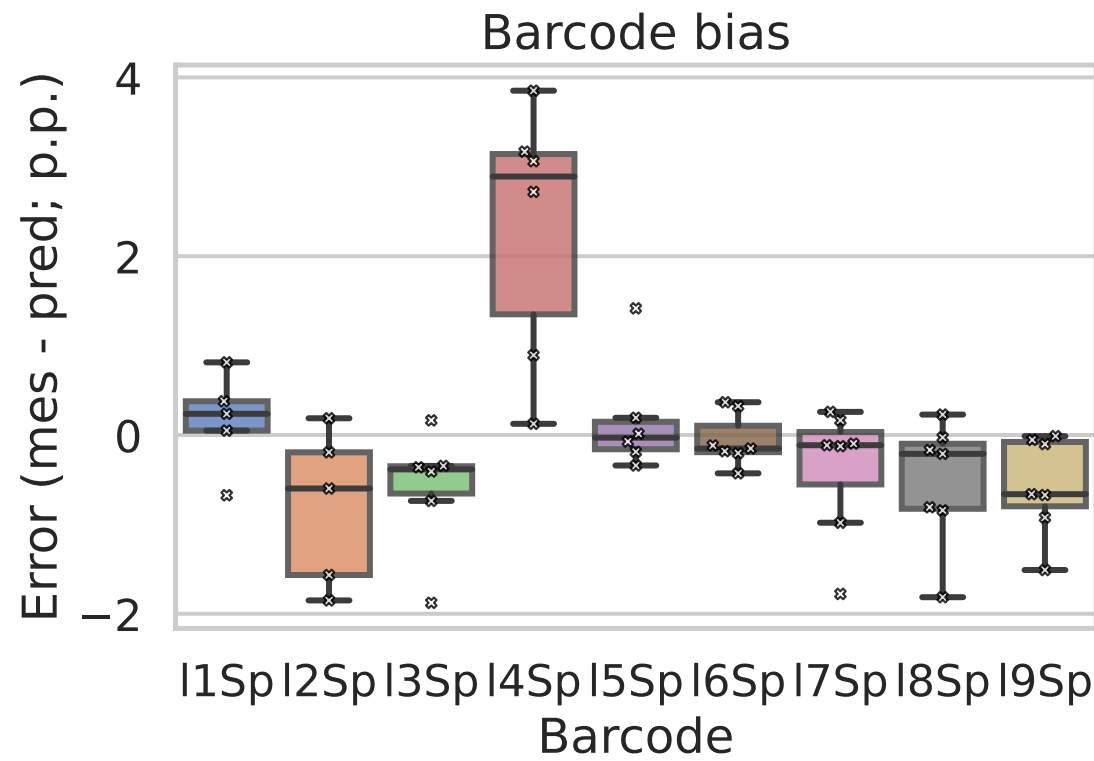
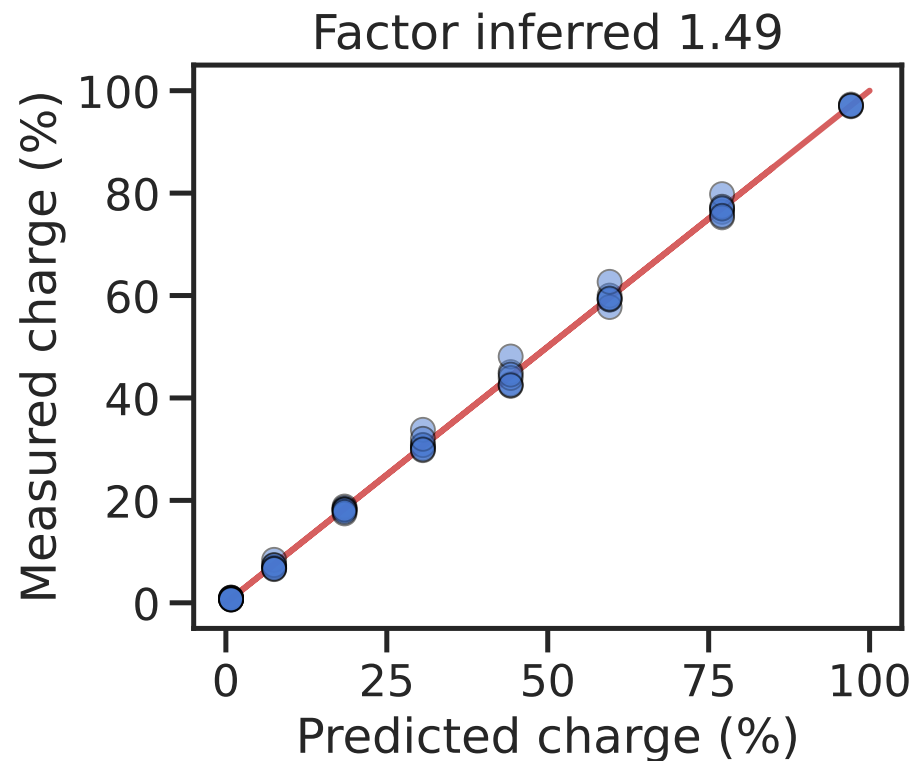
Leu-TAA-3-1



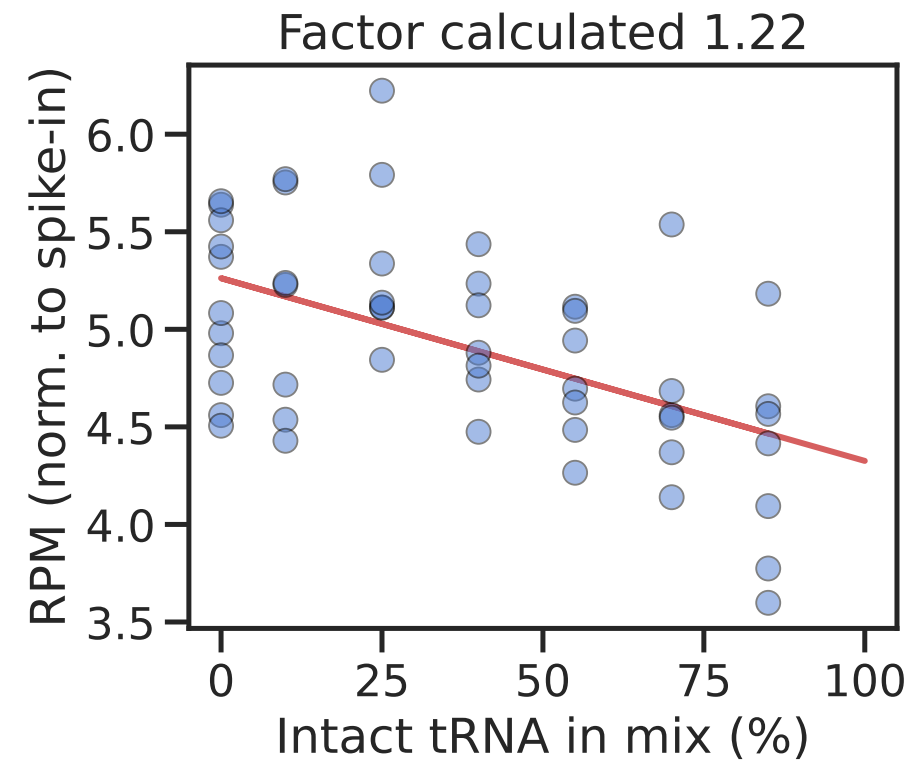
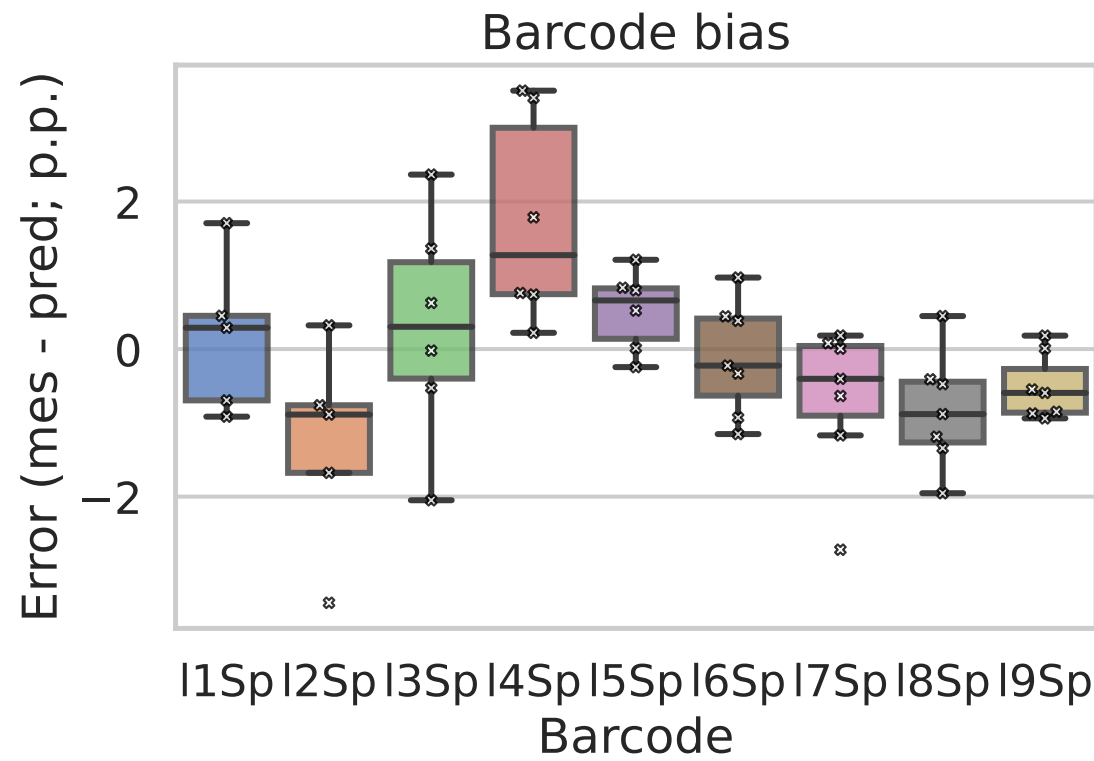
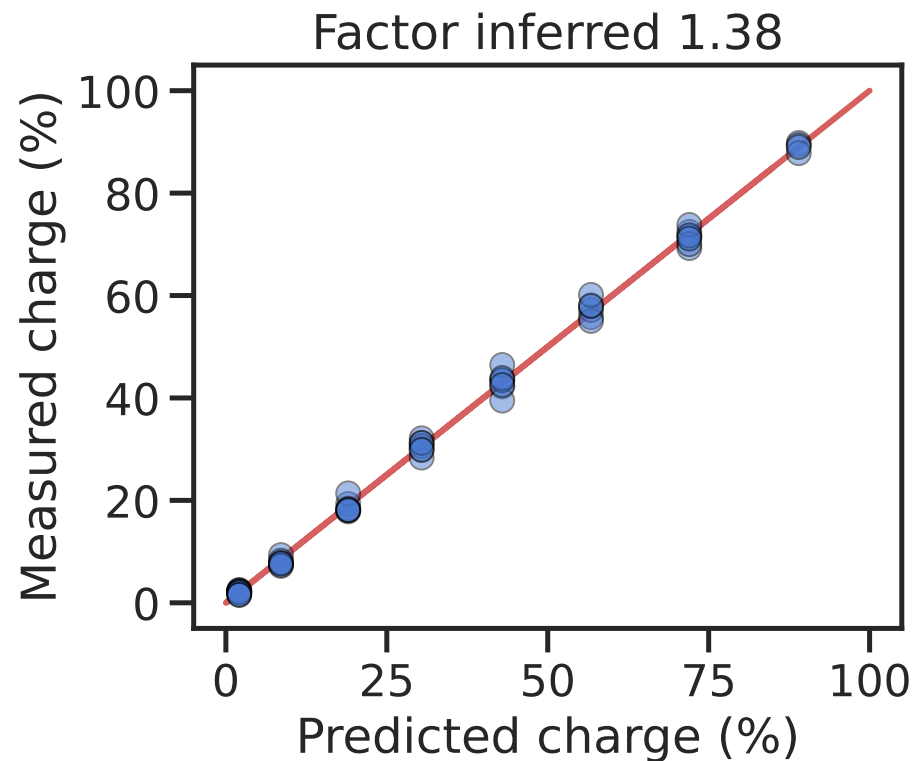
Ala-TGC-2-1



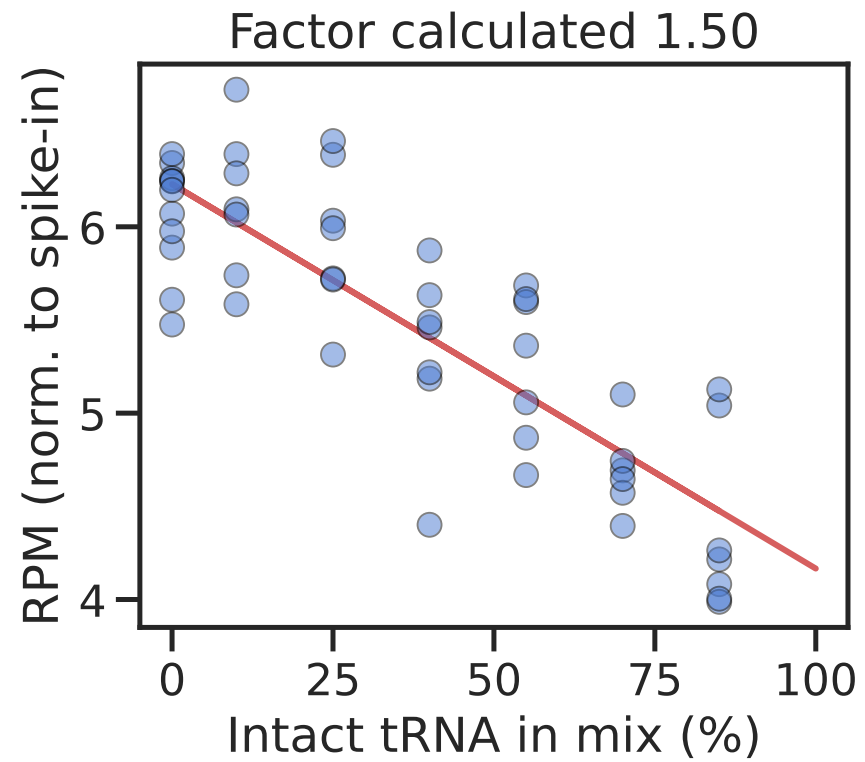
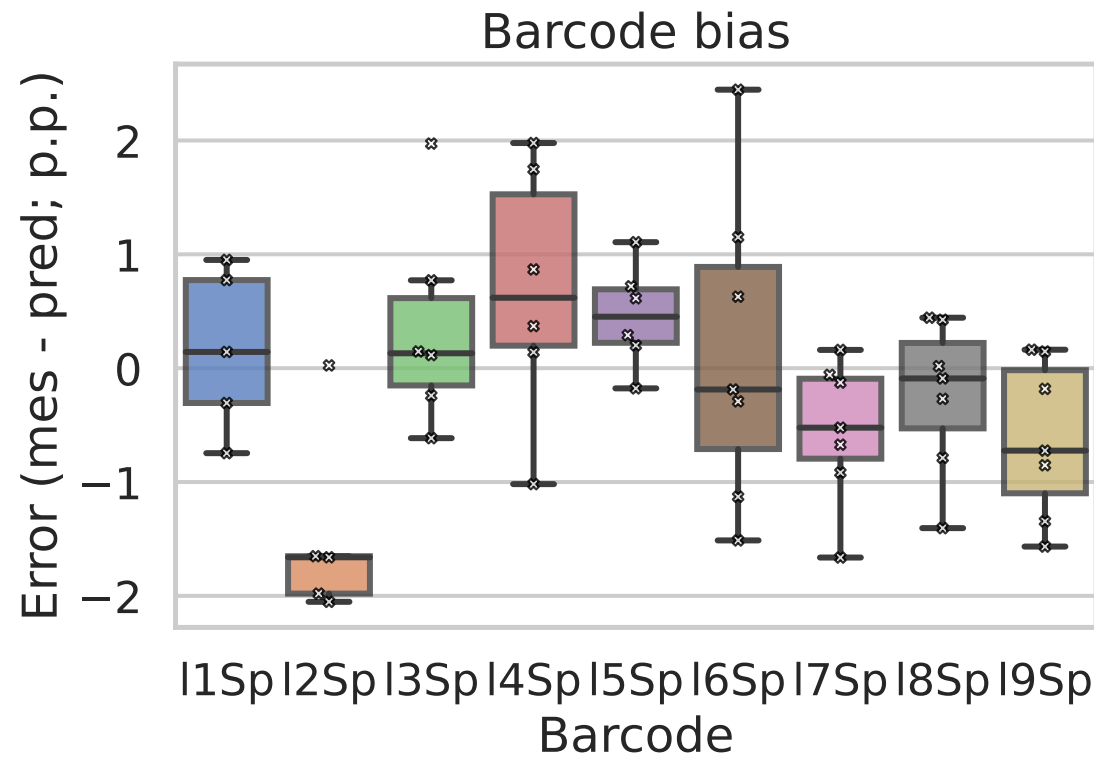
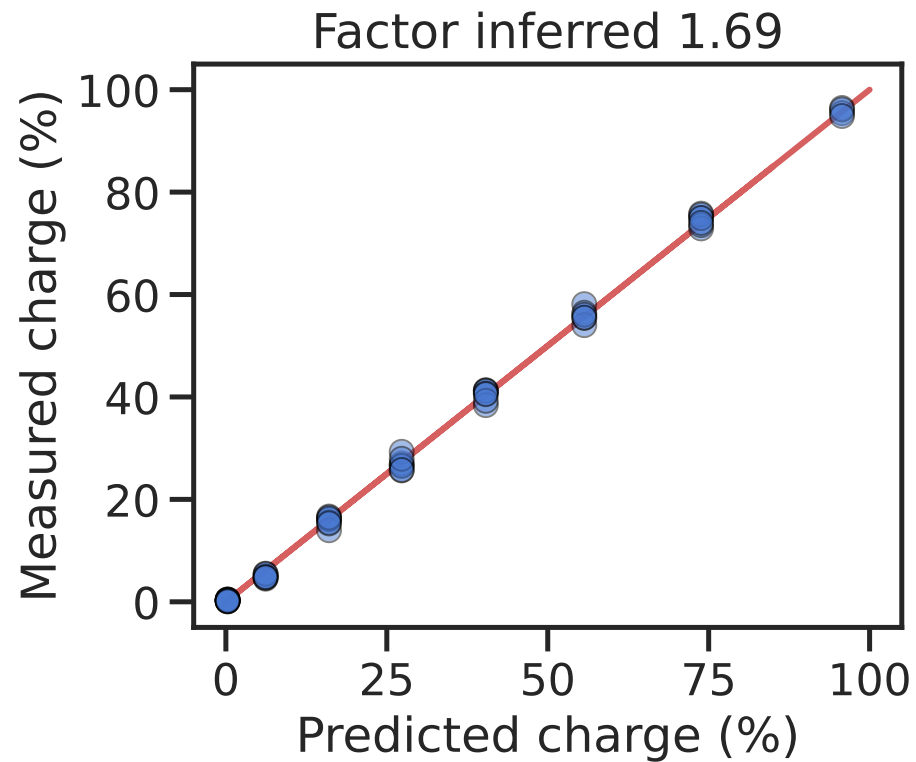
Gly-TCC-1-1



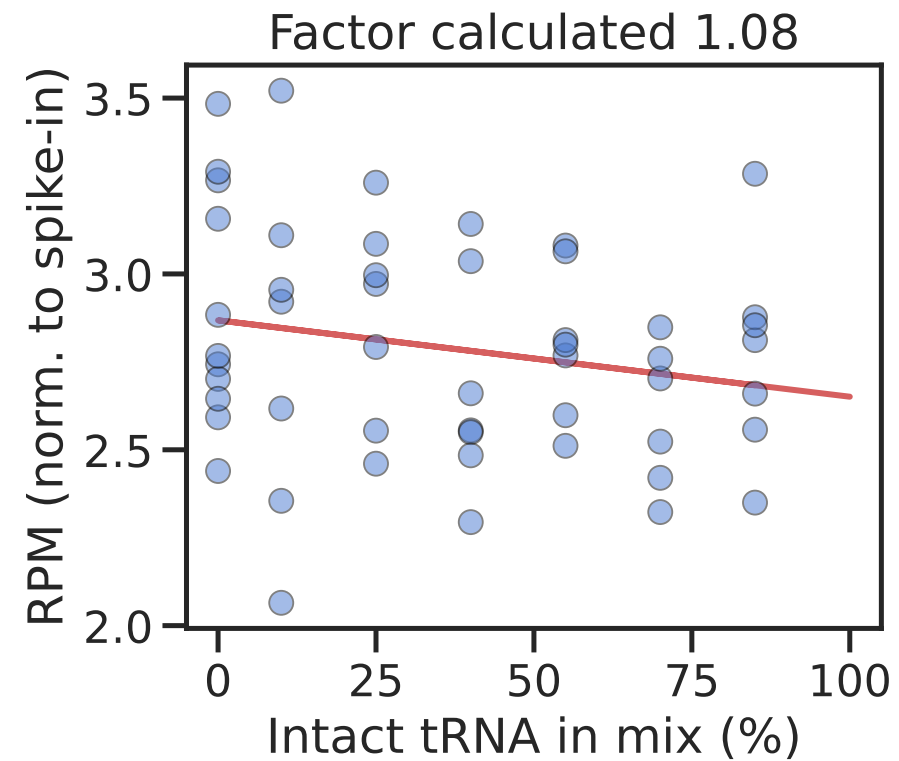
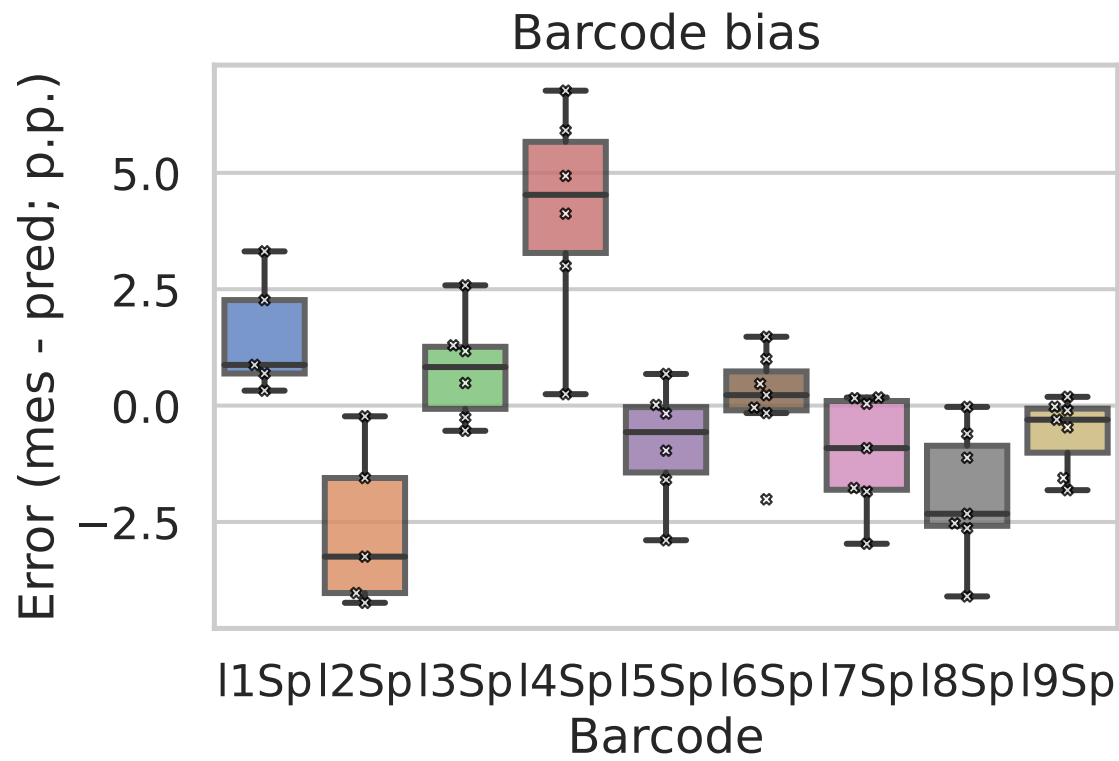
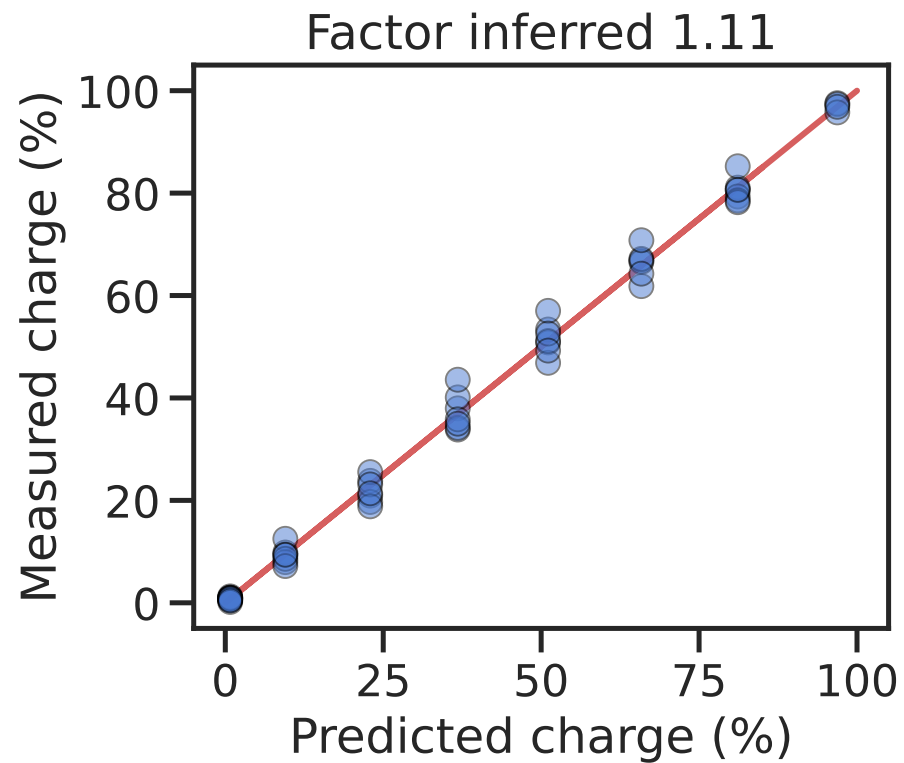
mito-Phe-GAA



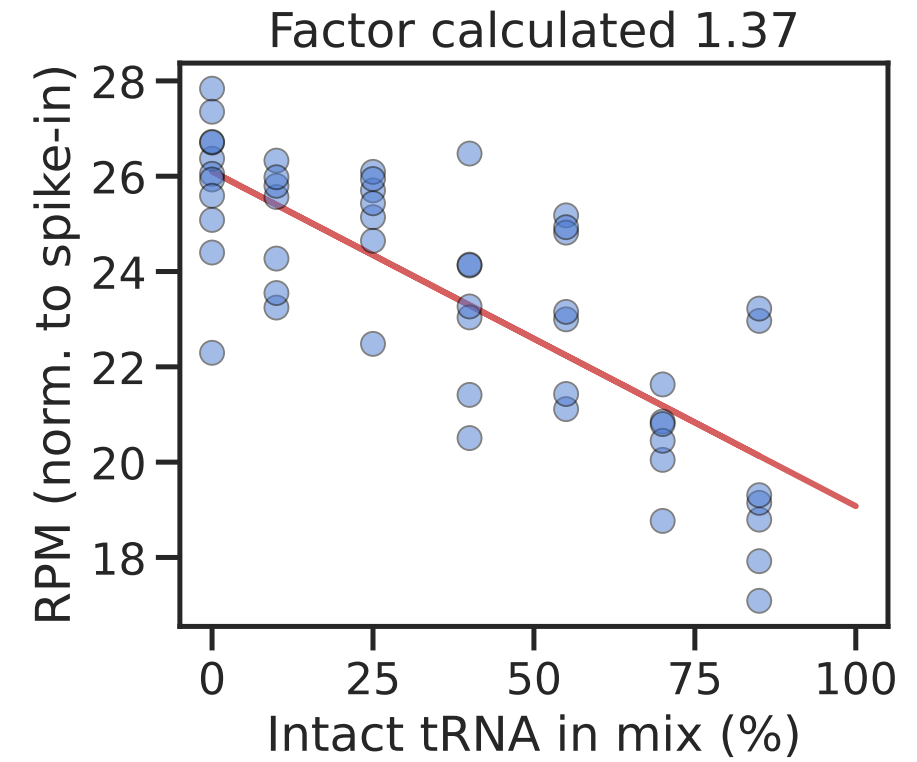
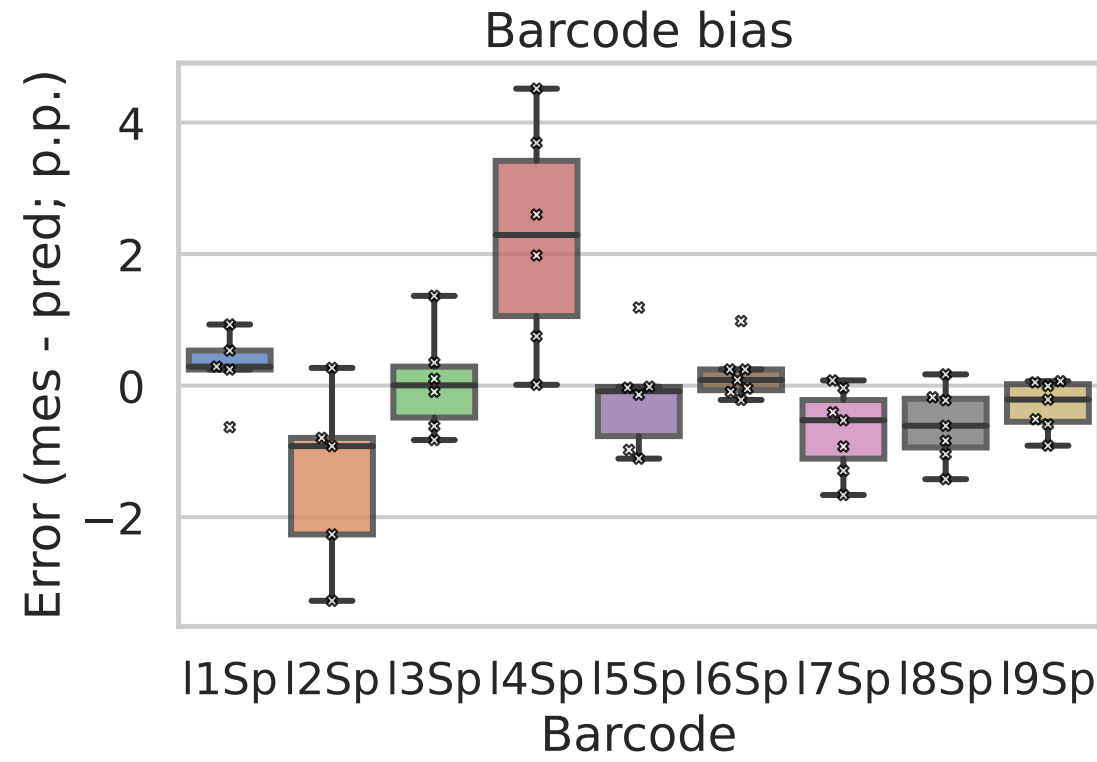
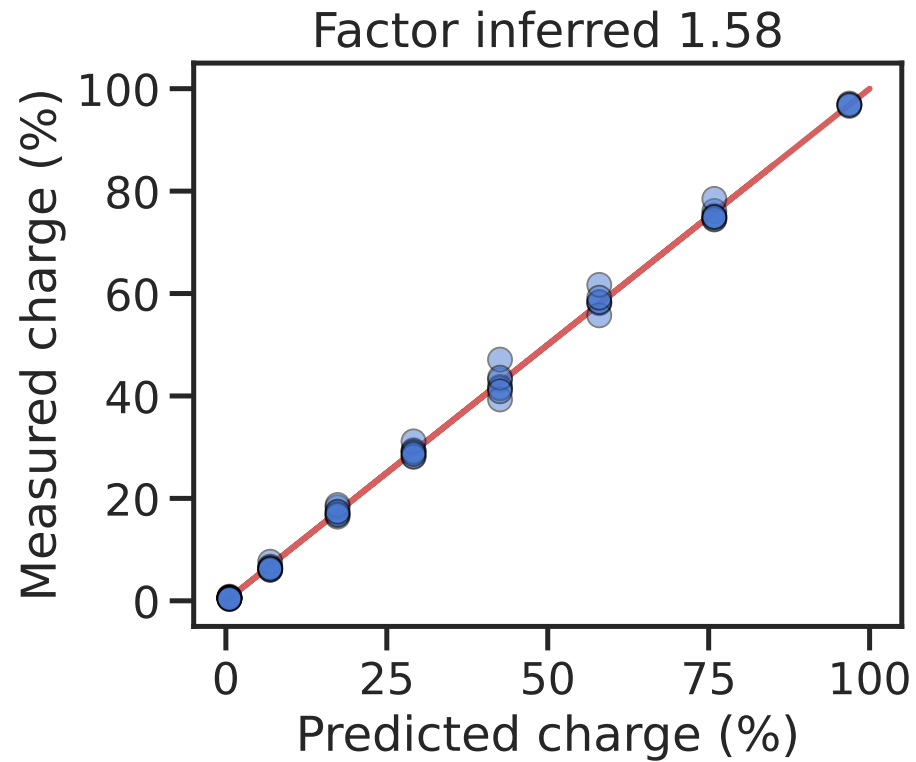
Asn-GTT-5-1



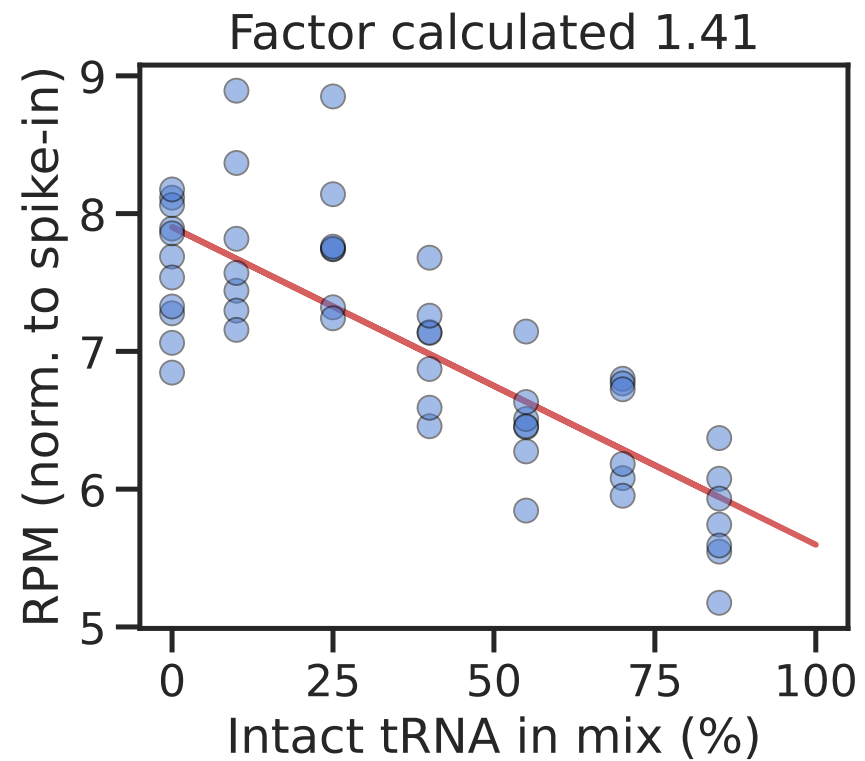
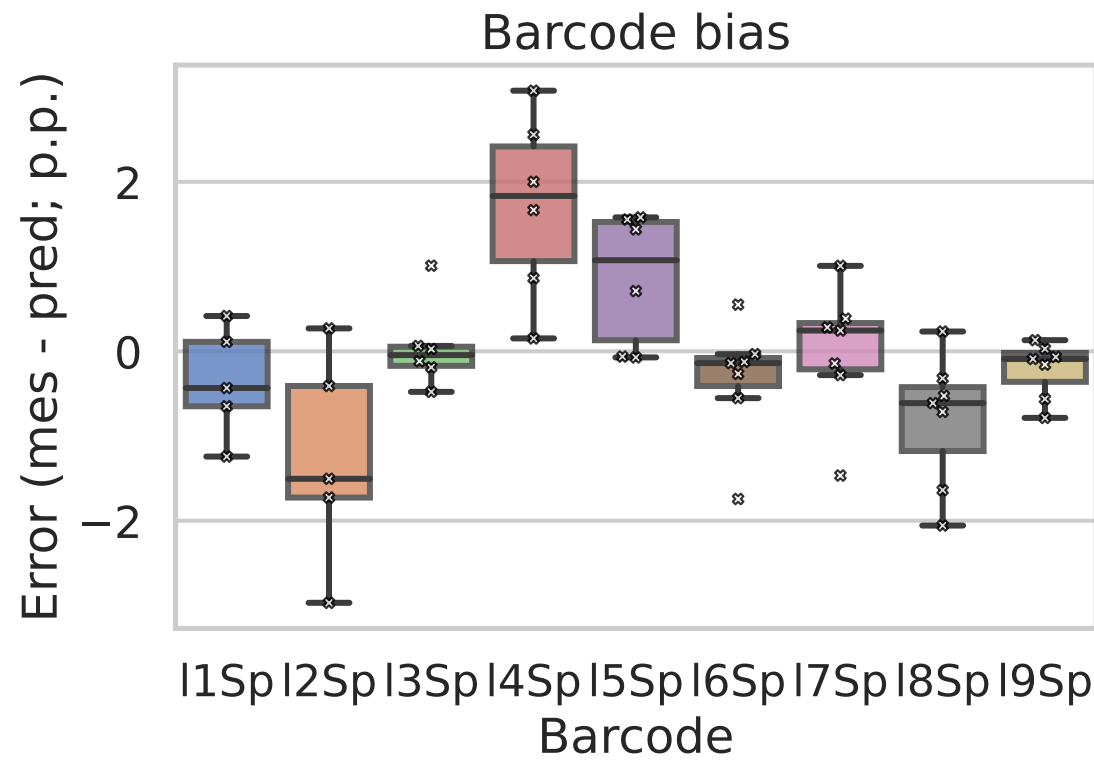
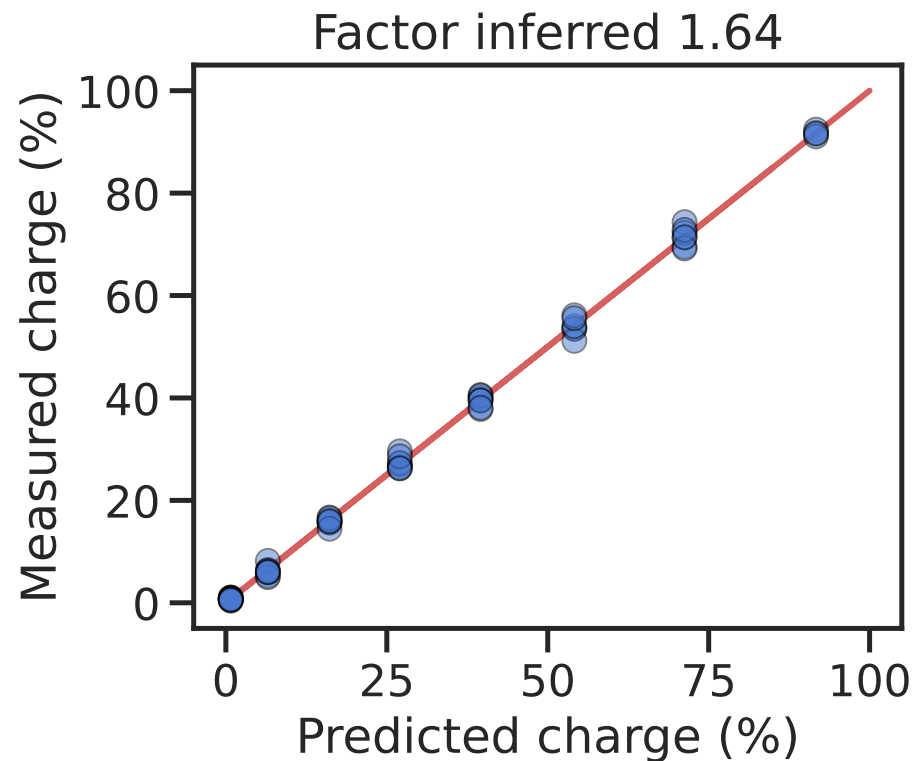
Cys-GCA-9-1



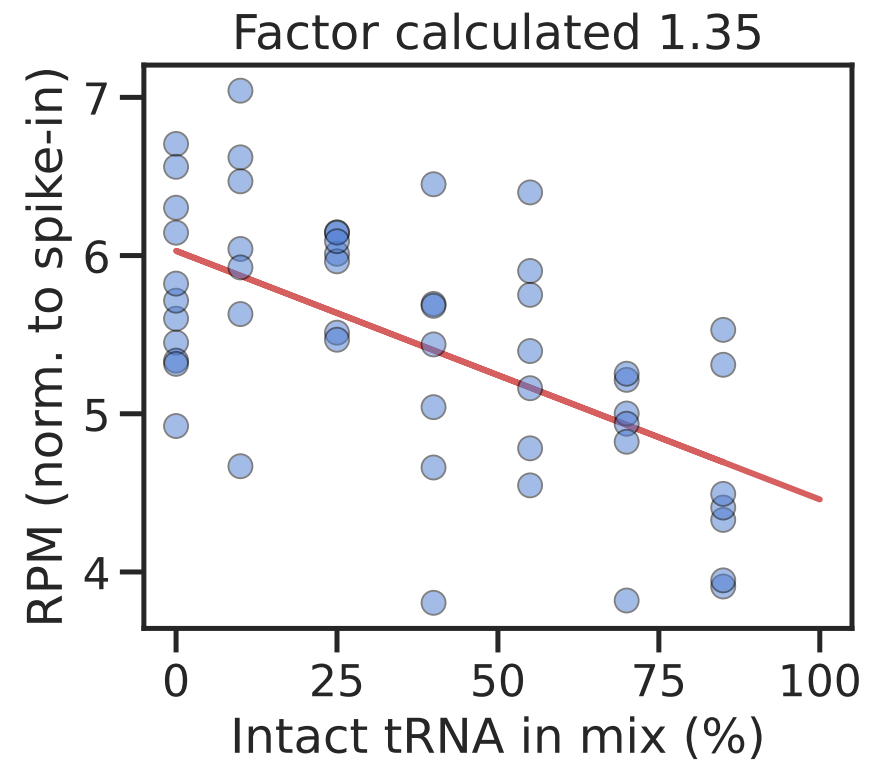
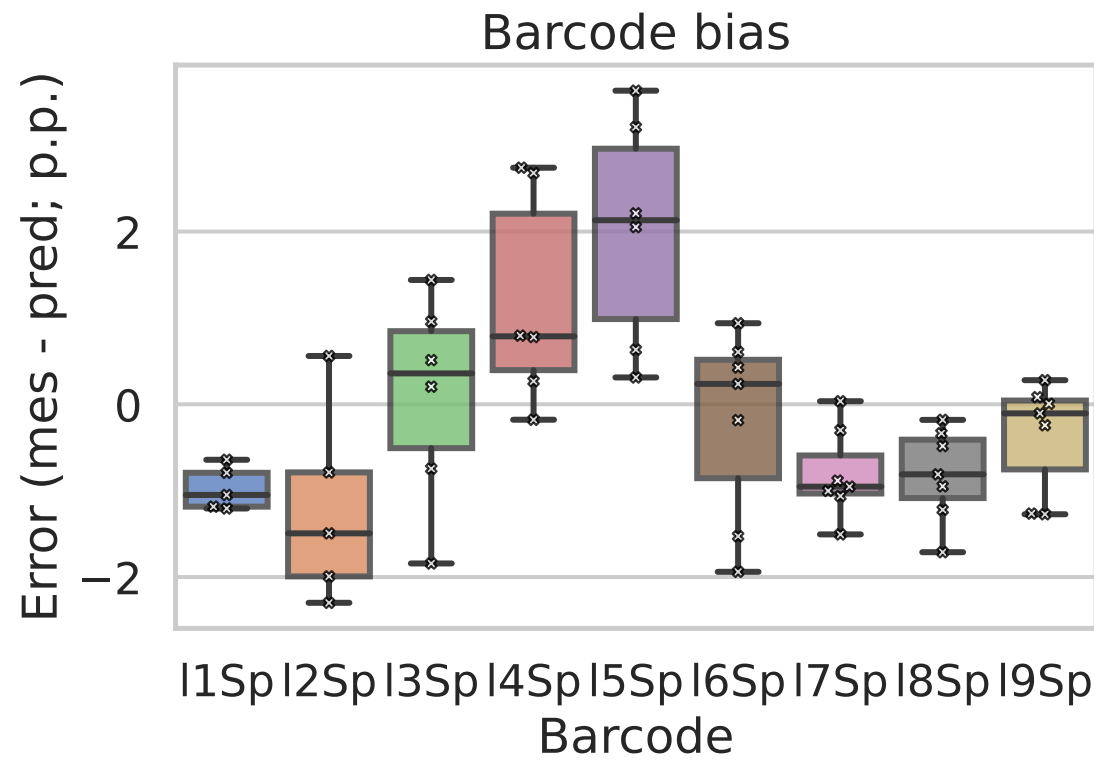
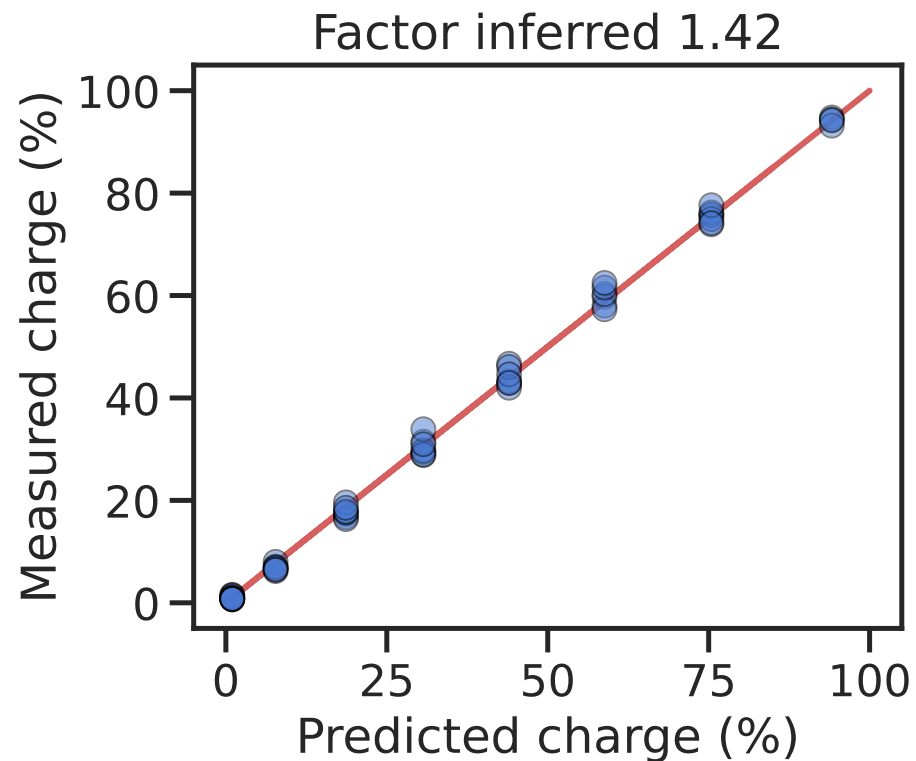
Ala-TGC-3-2



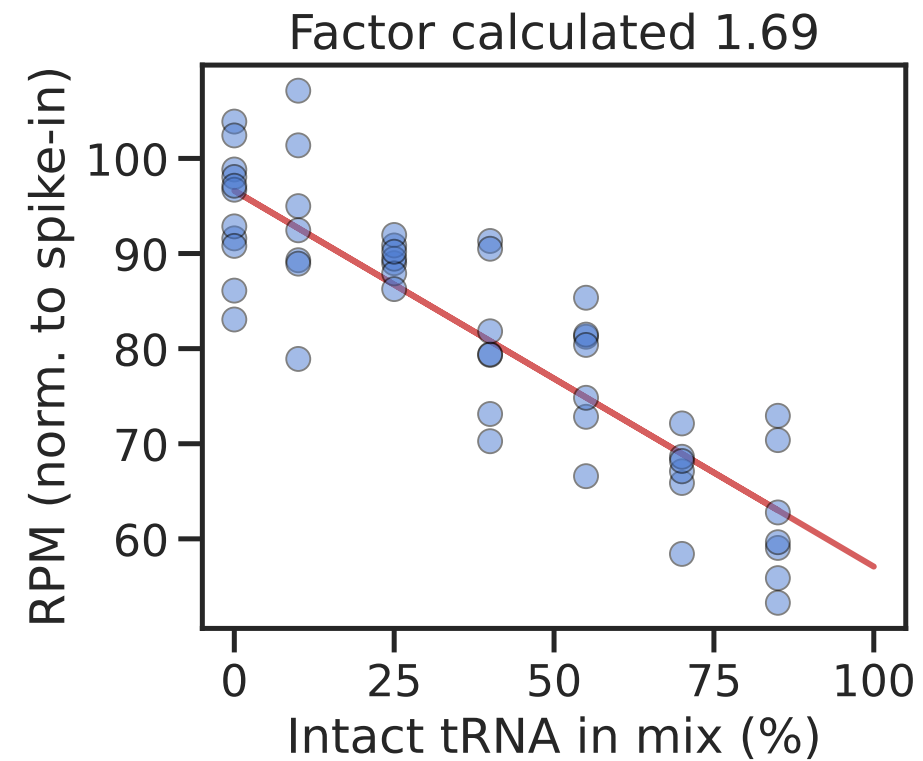
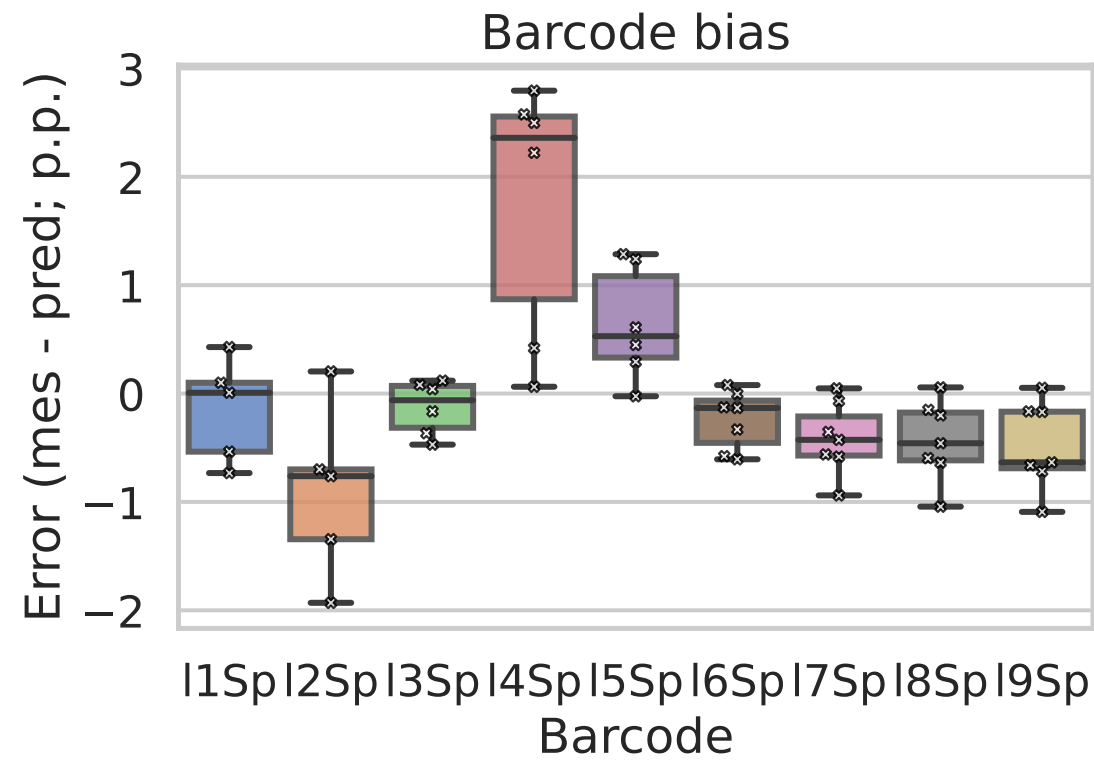
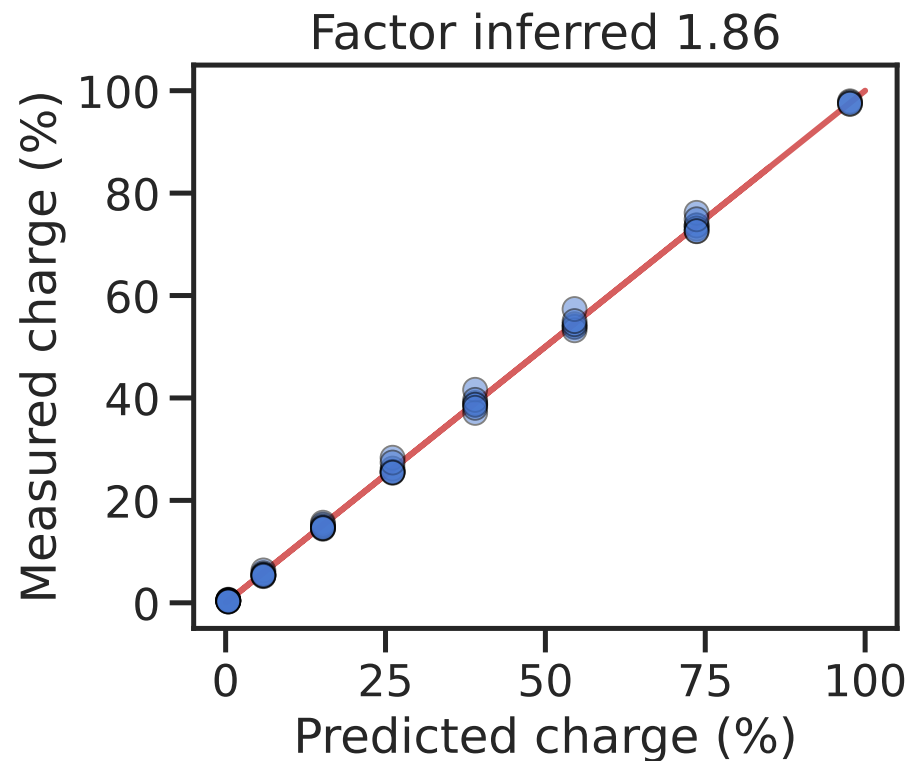
mito-Glu-TTC



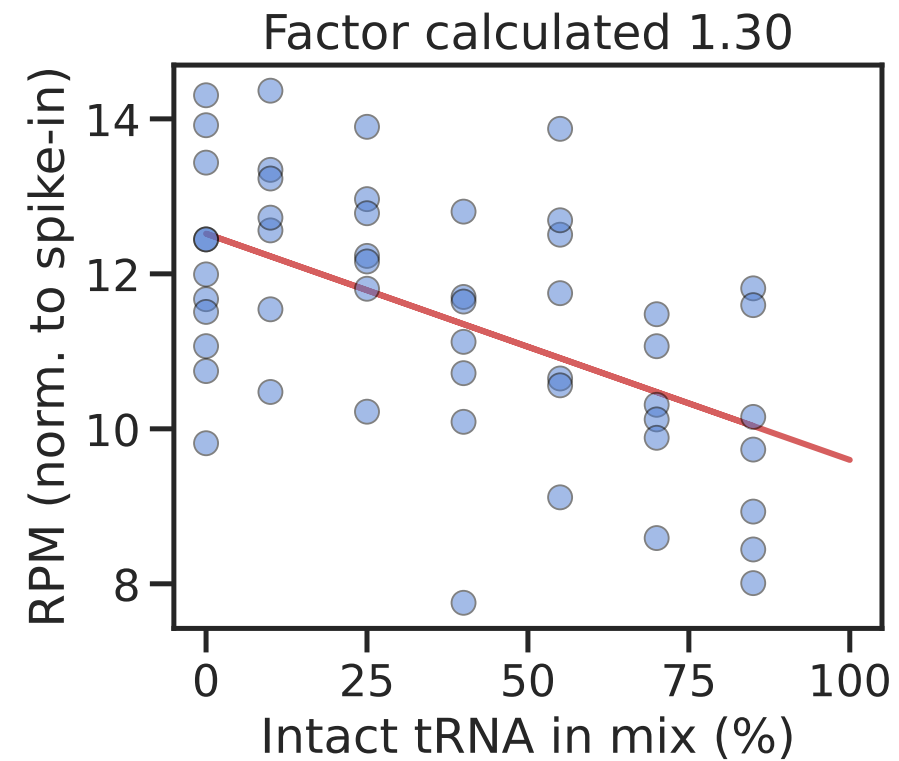
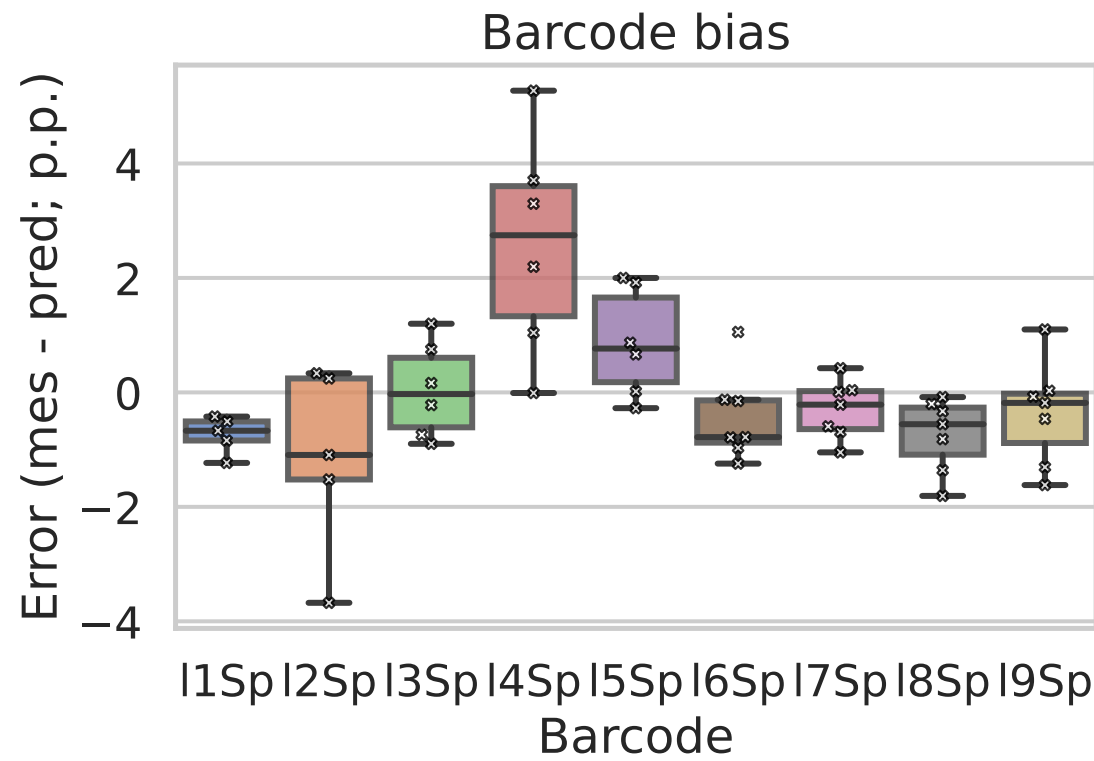
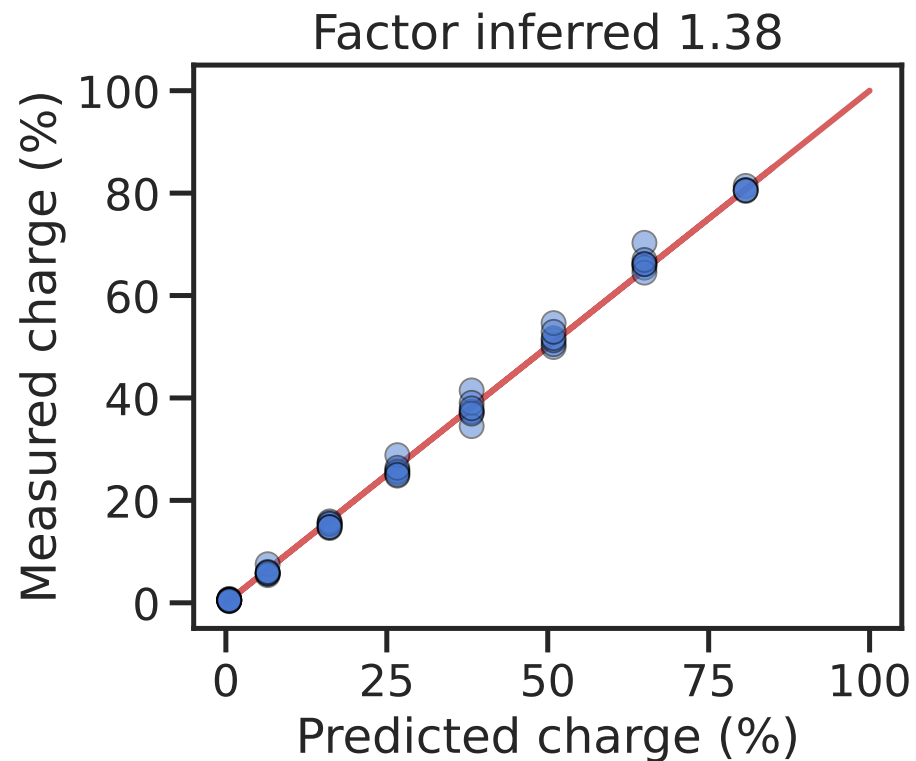
Asp-GTC-1-1



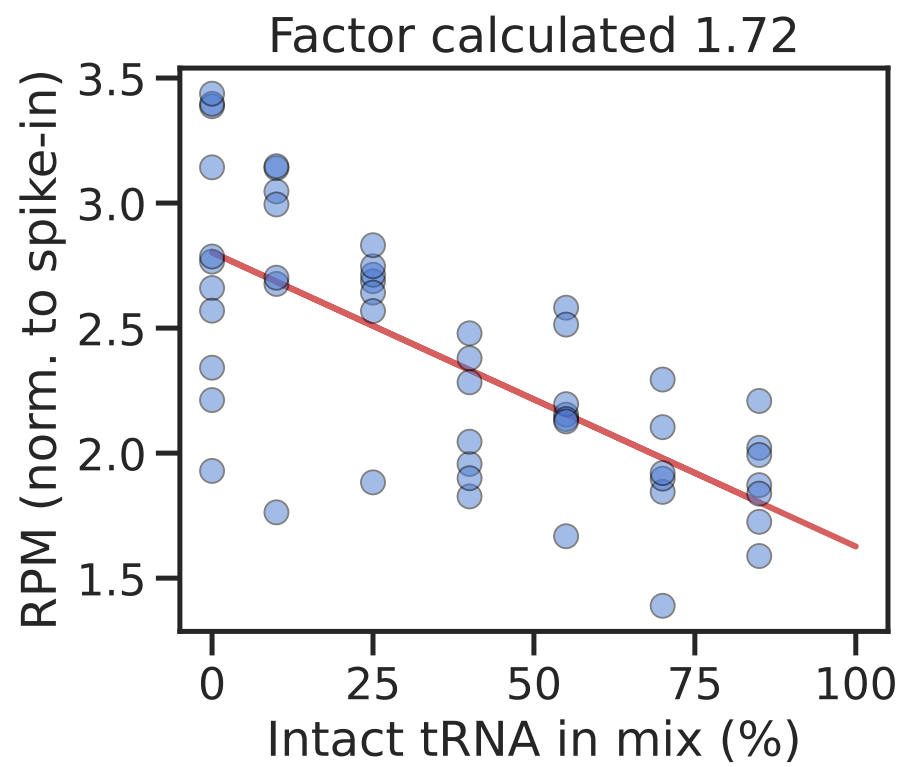
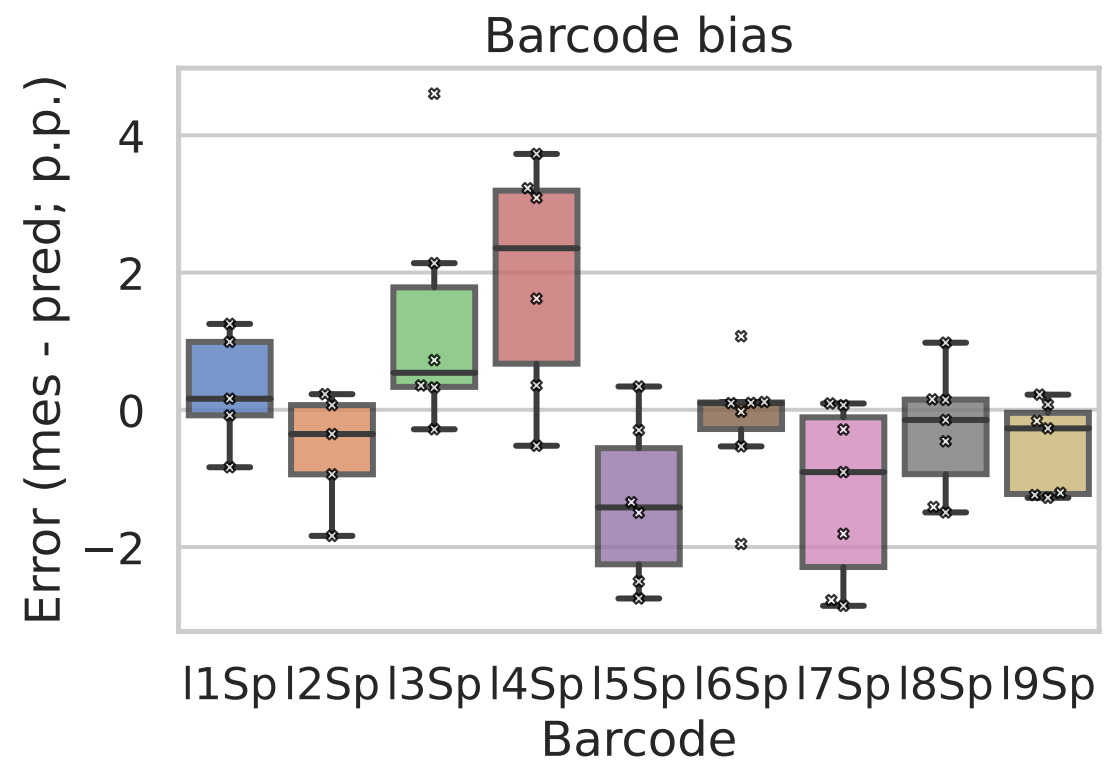
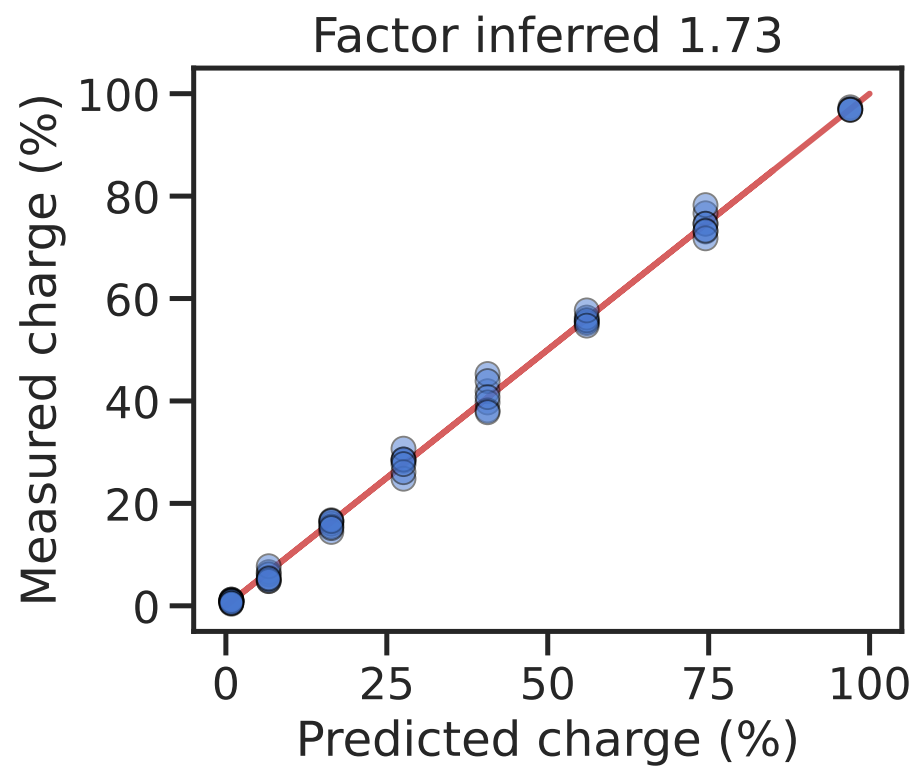
iMet-CAT-1-1



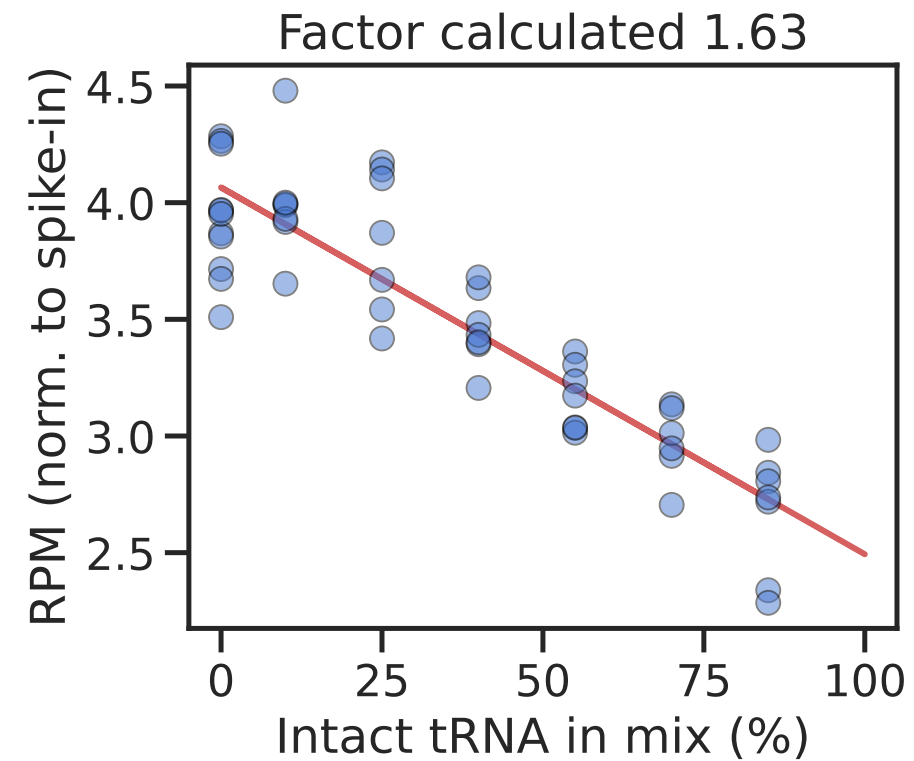
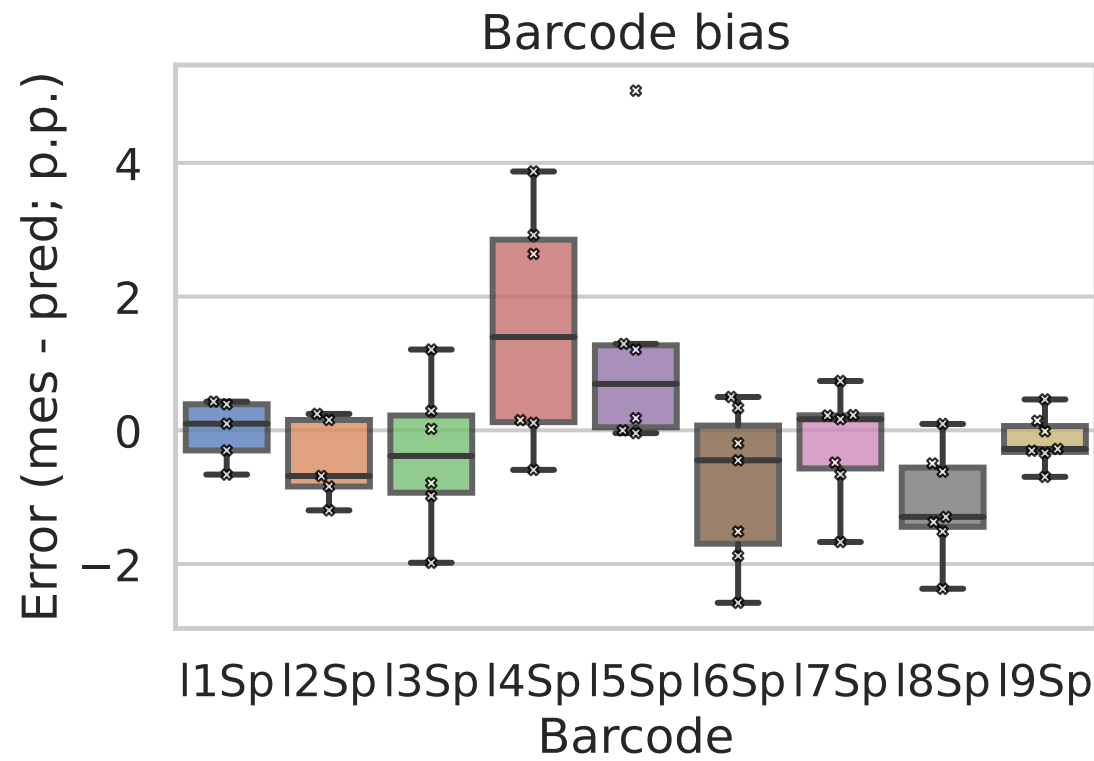
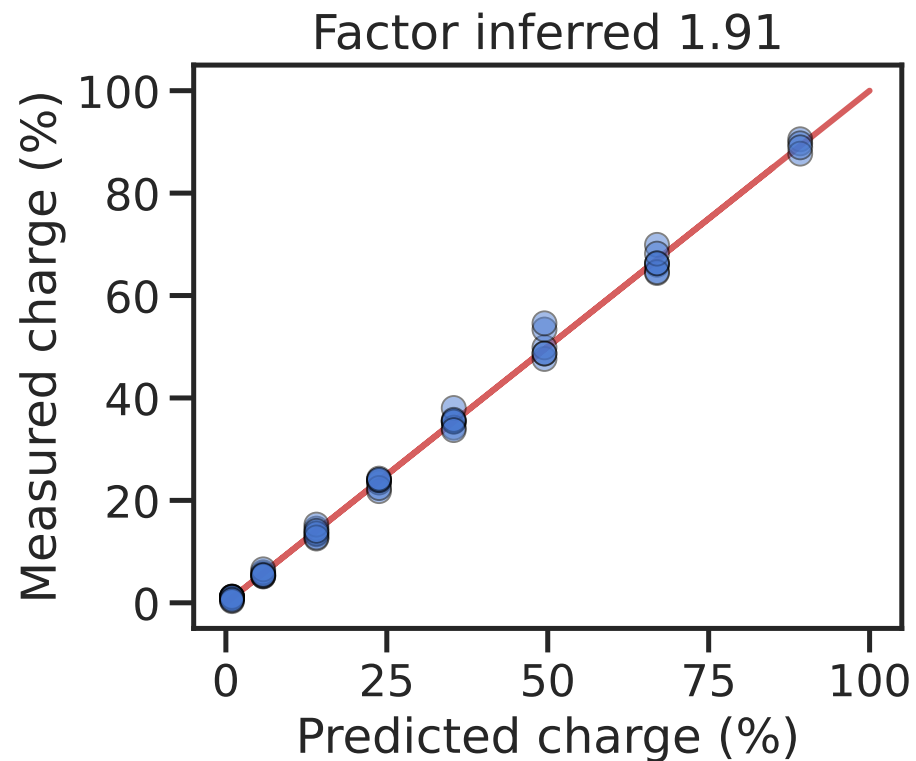
Glu-TTC-4-1



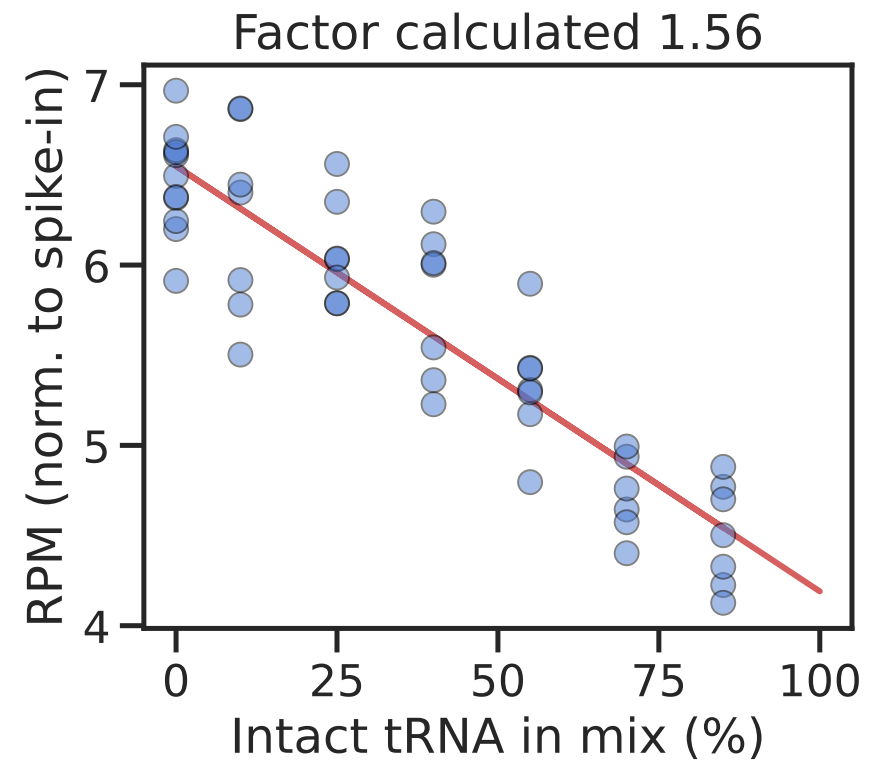
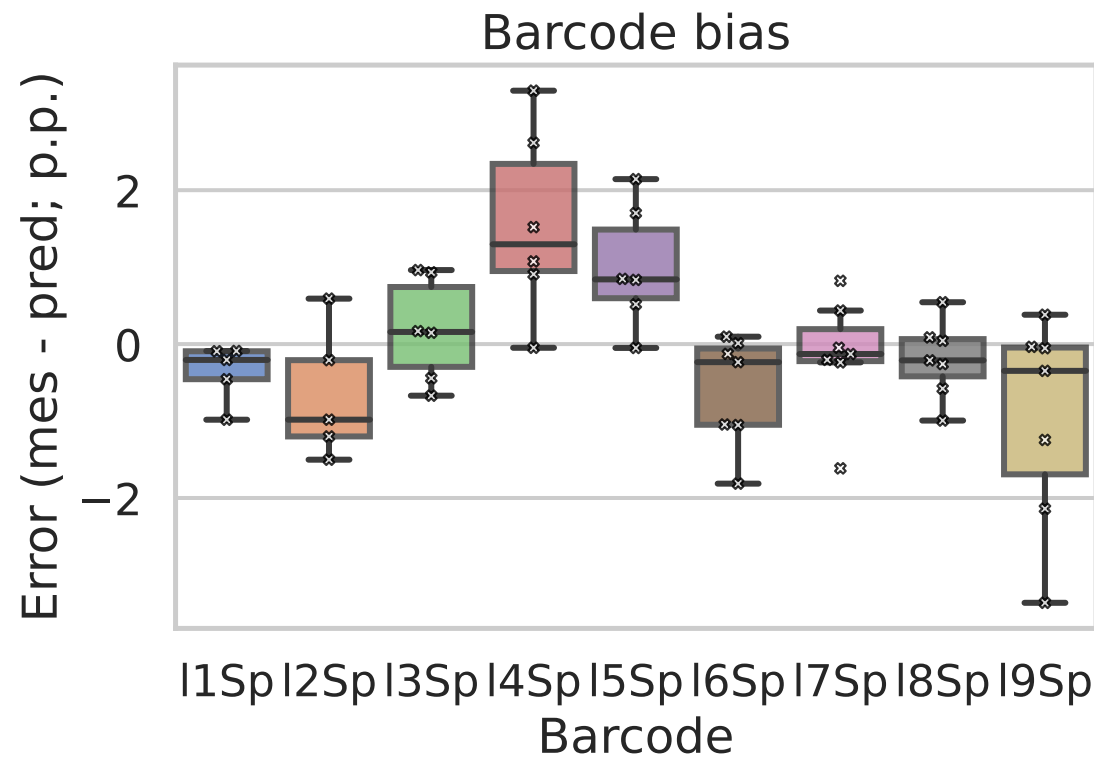
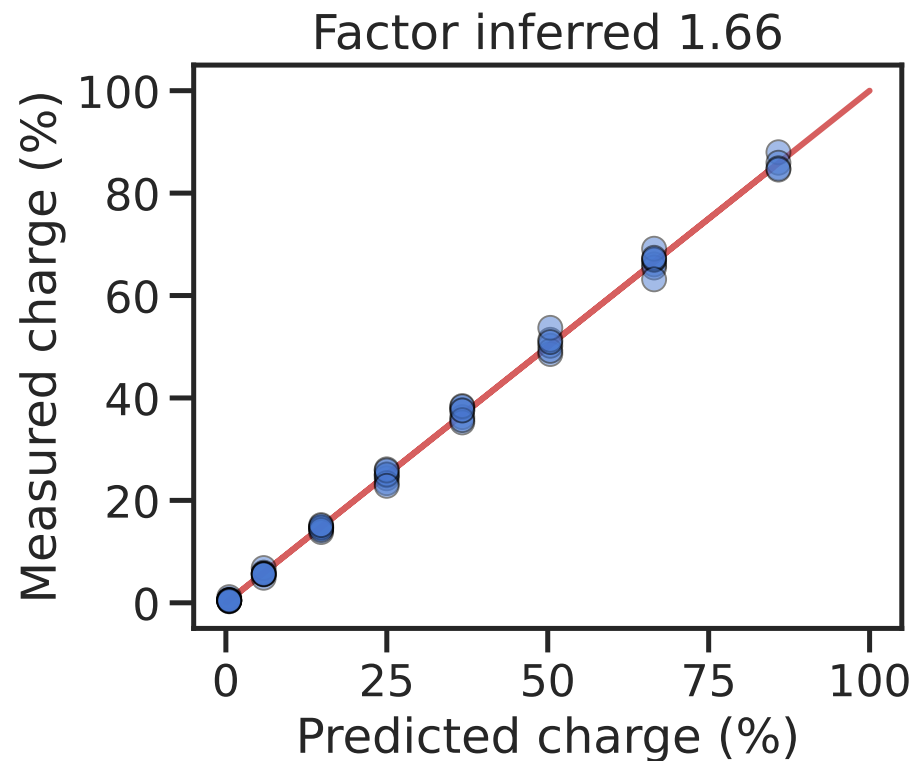
Trp-CCA-1-1



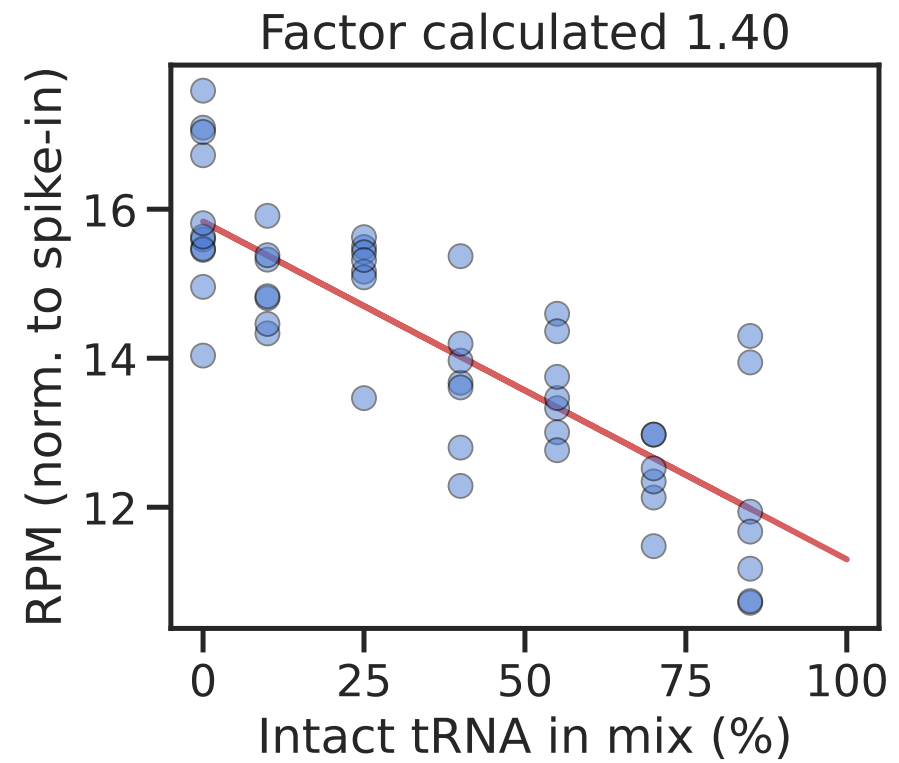
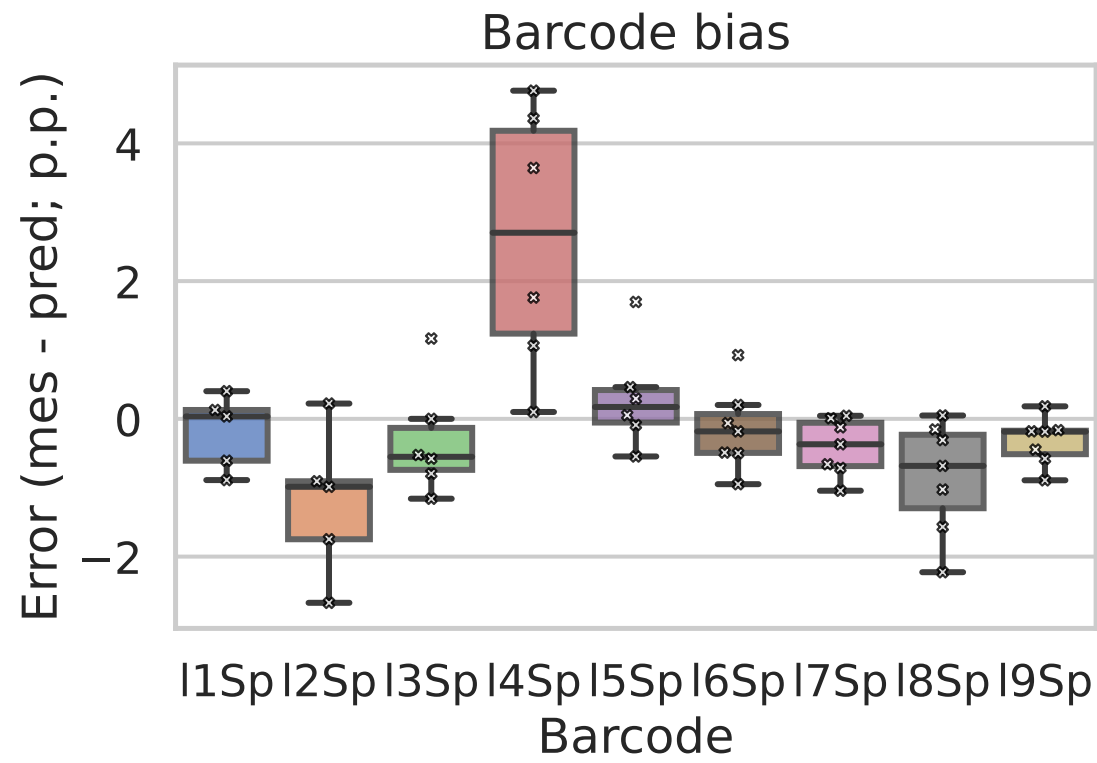
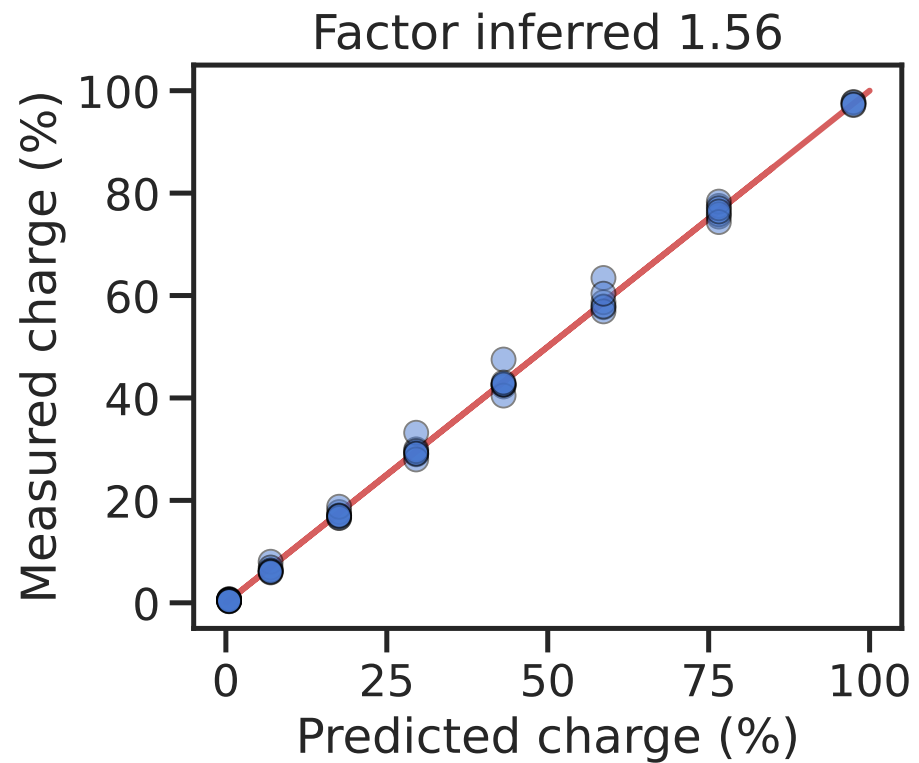
mito-Leu1-TAG



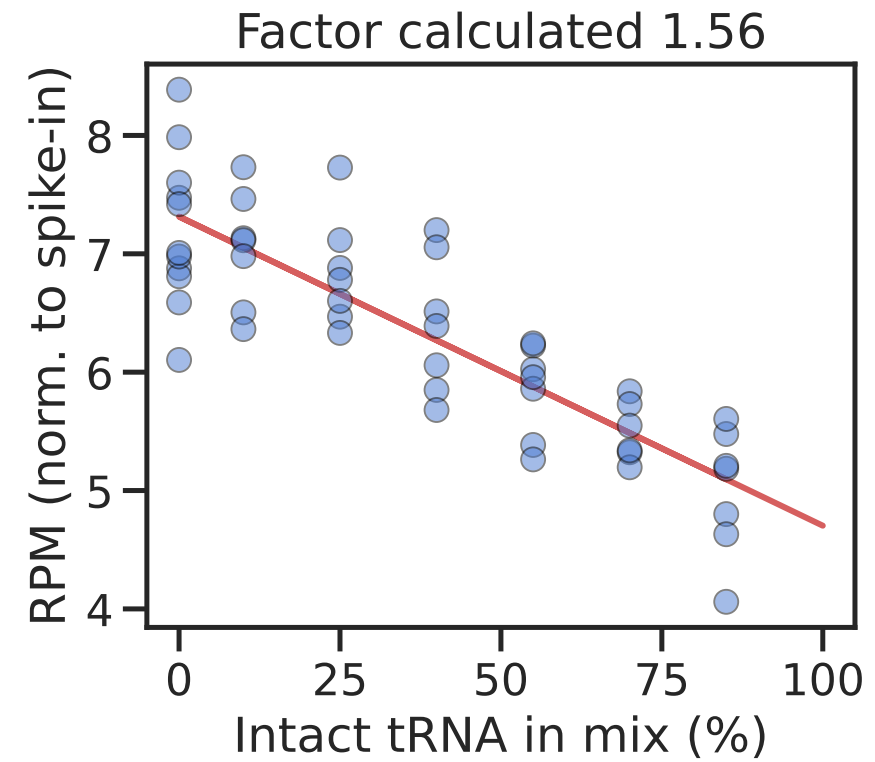
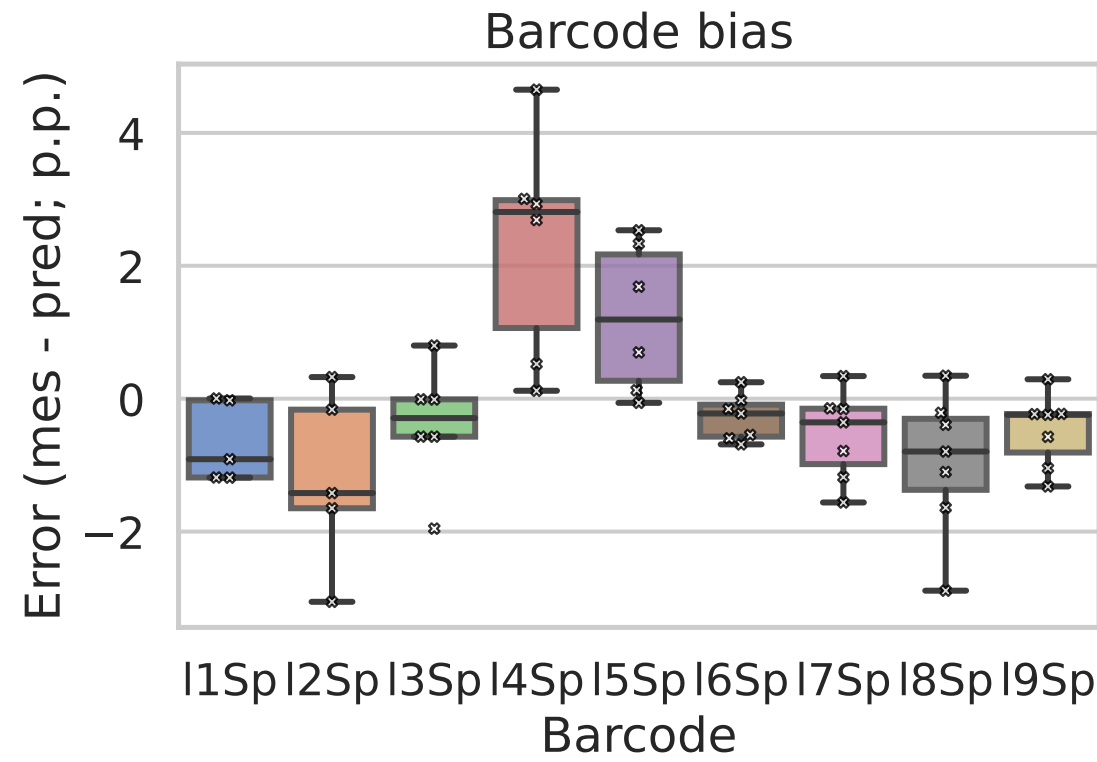
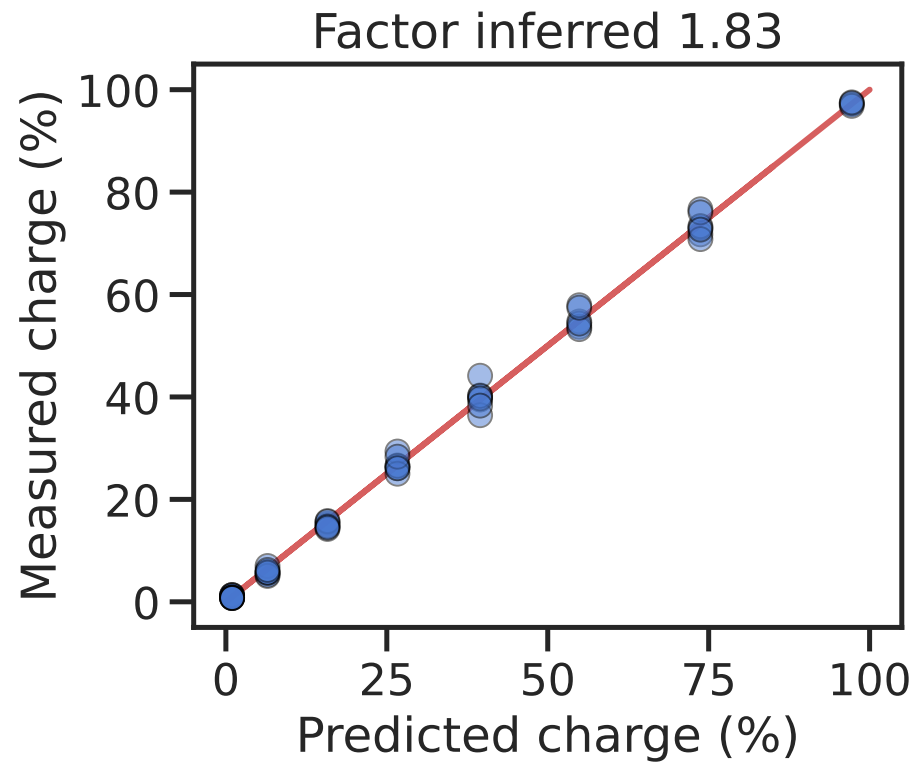
mito-His-GTG



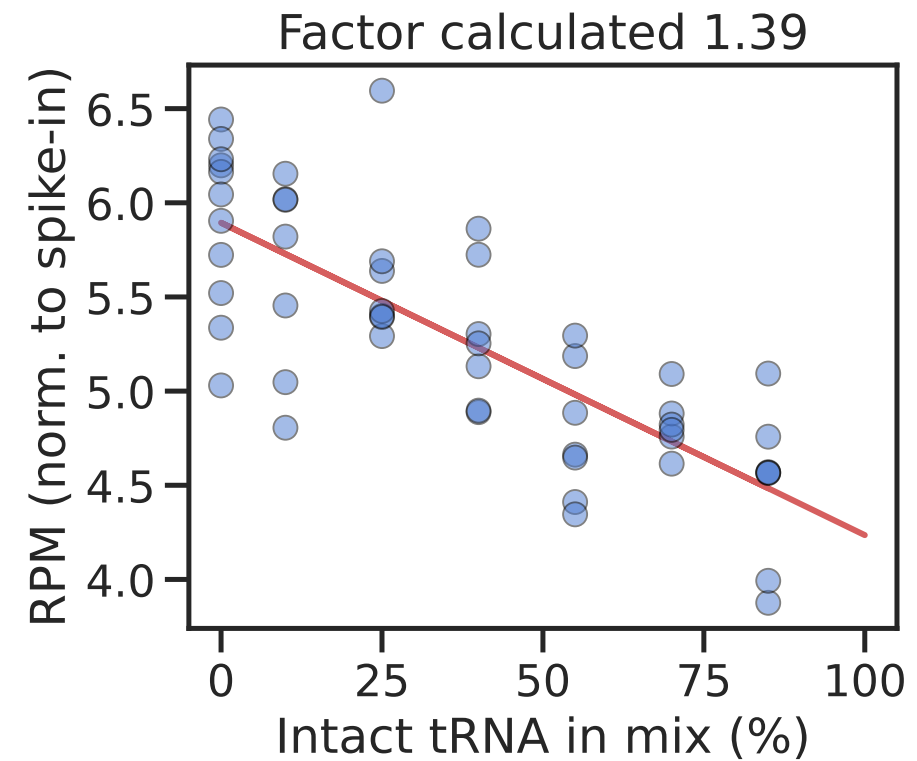
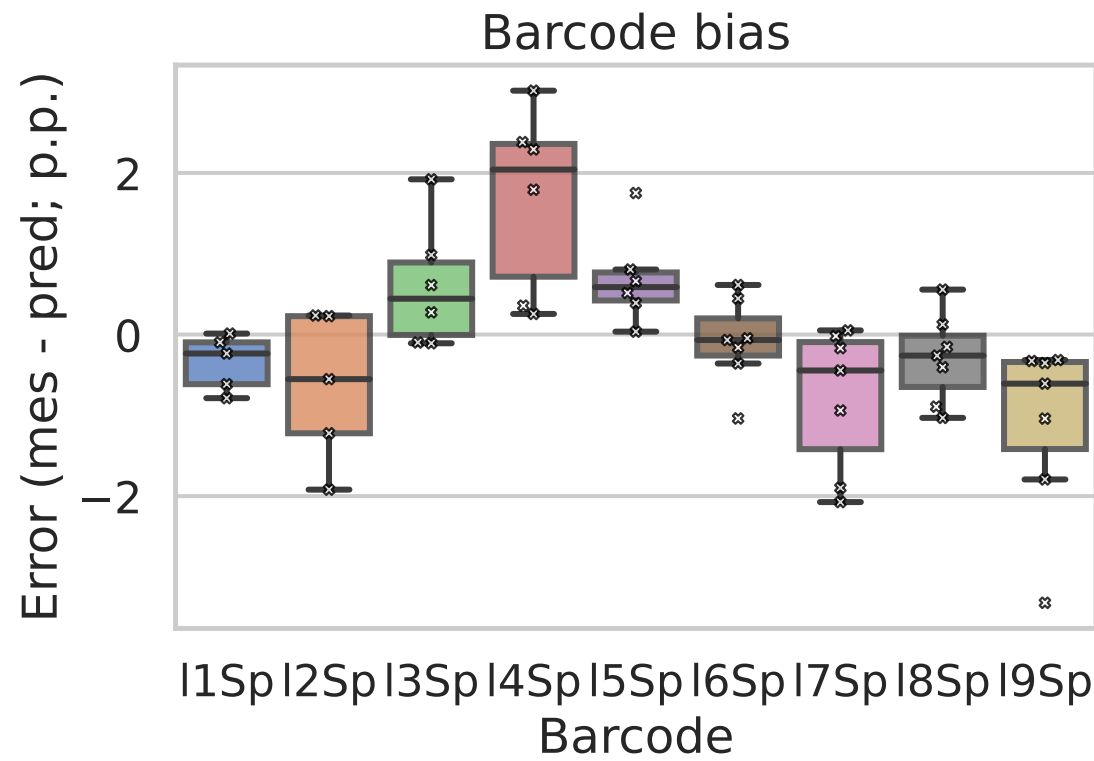
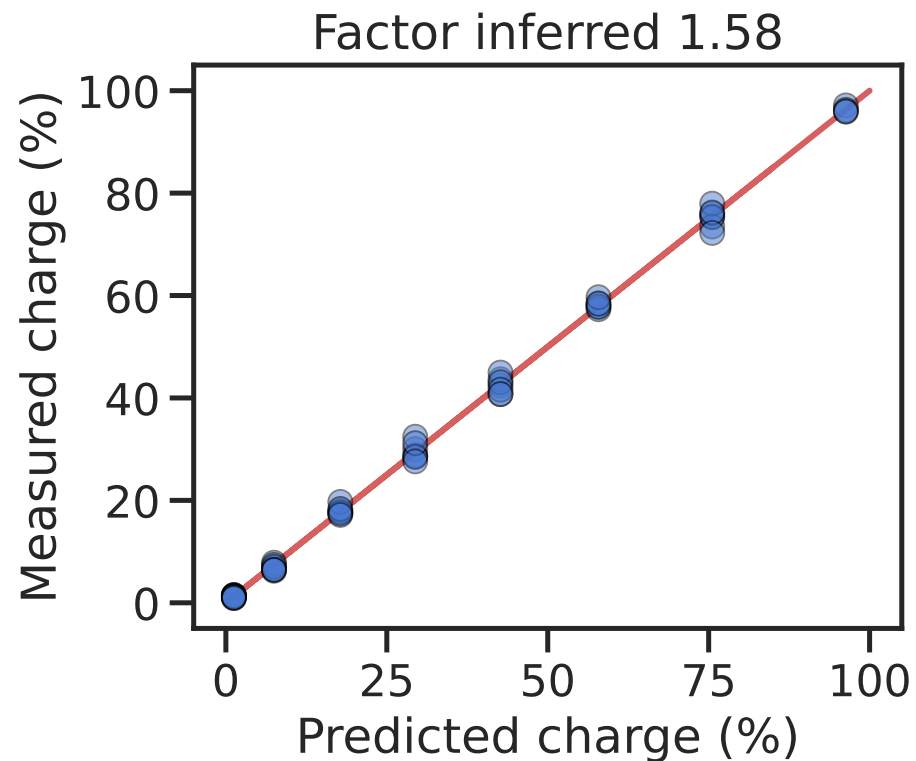
Ala-CGC-1-1



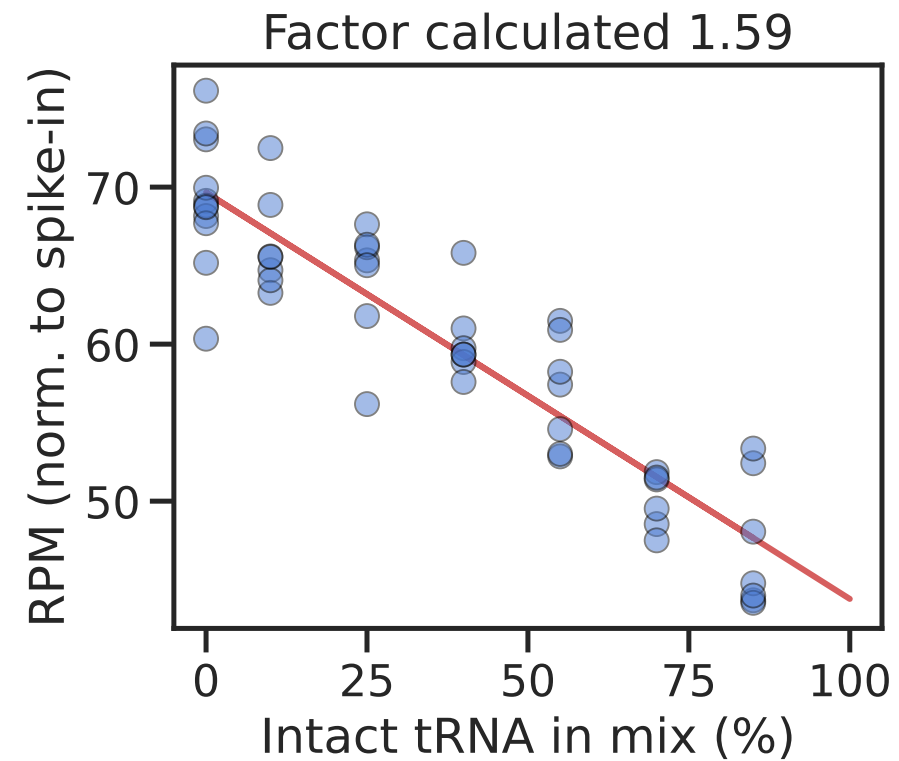
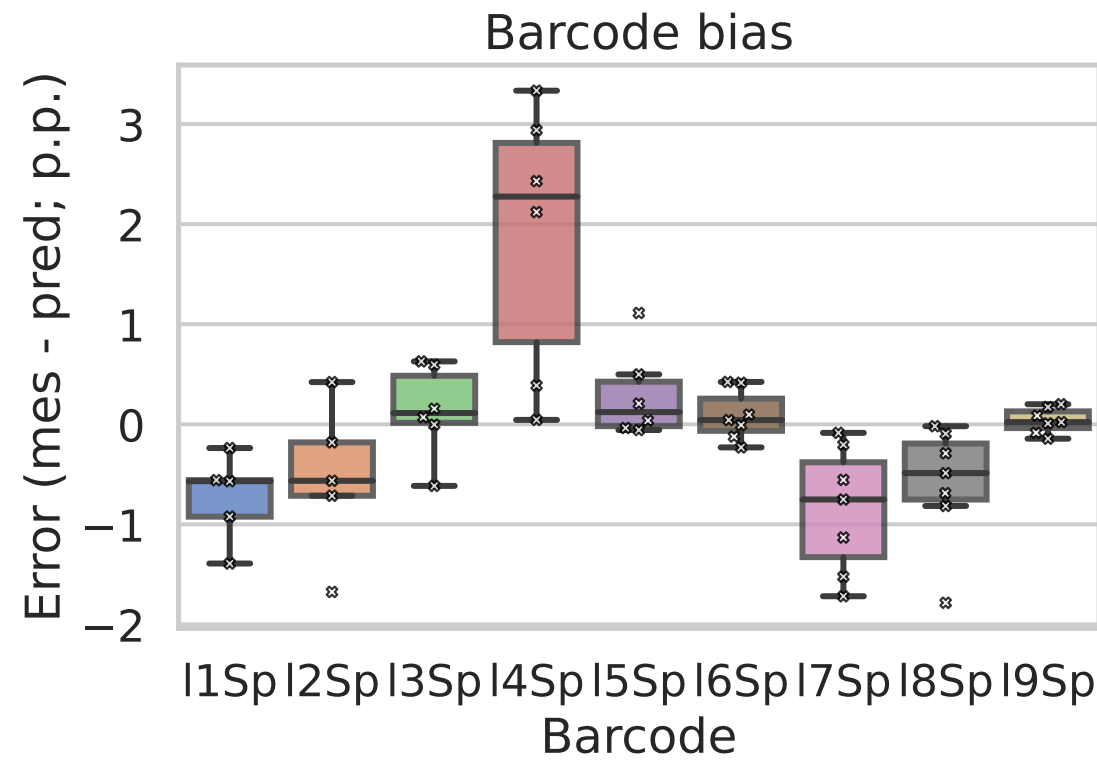
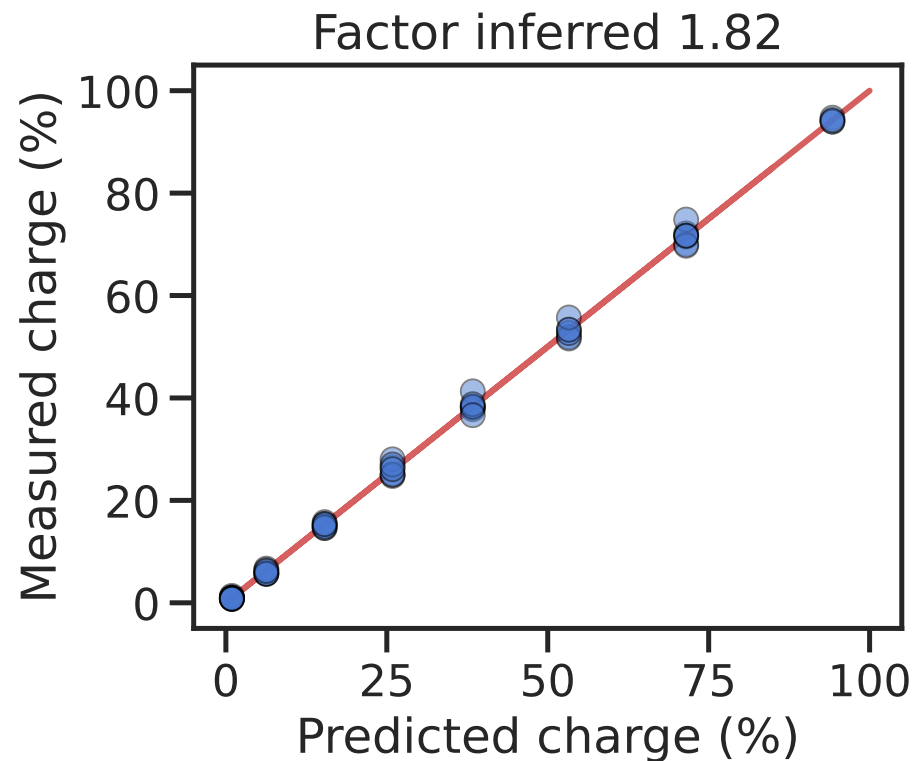
mito-Leu2-TAA



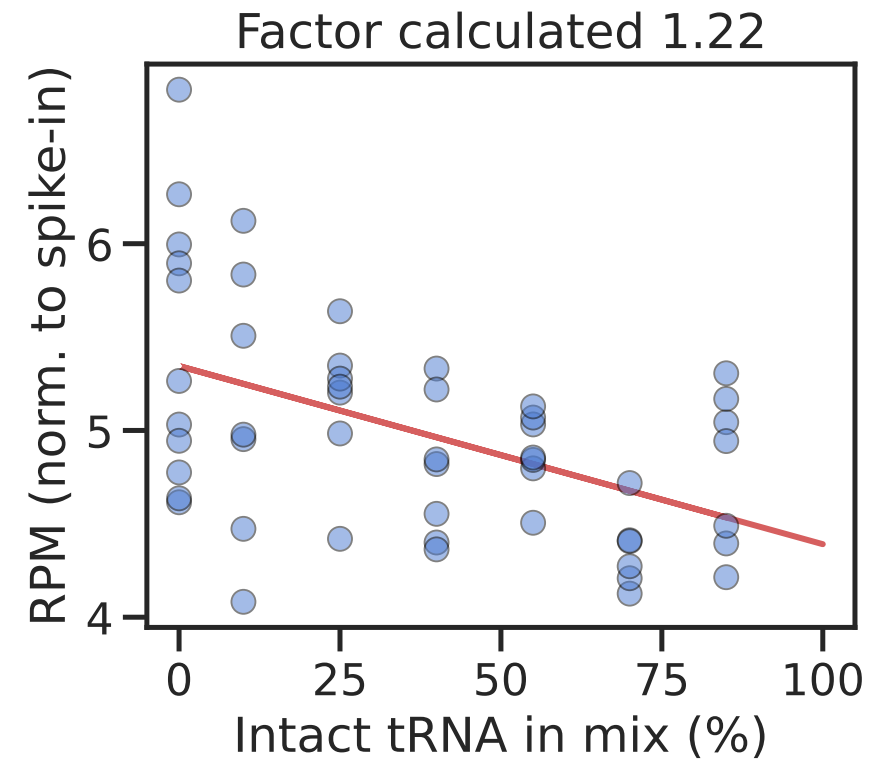
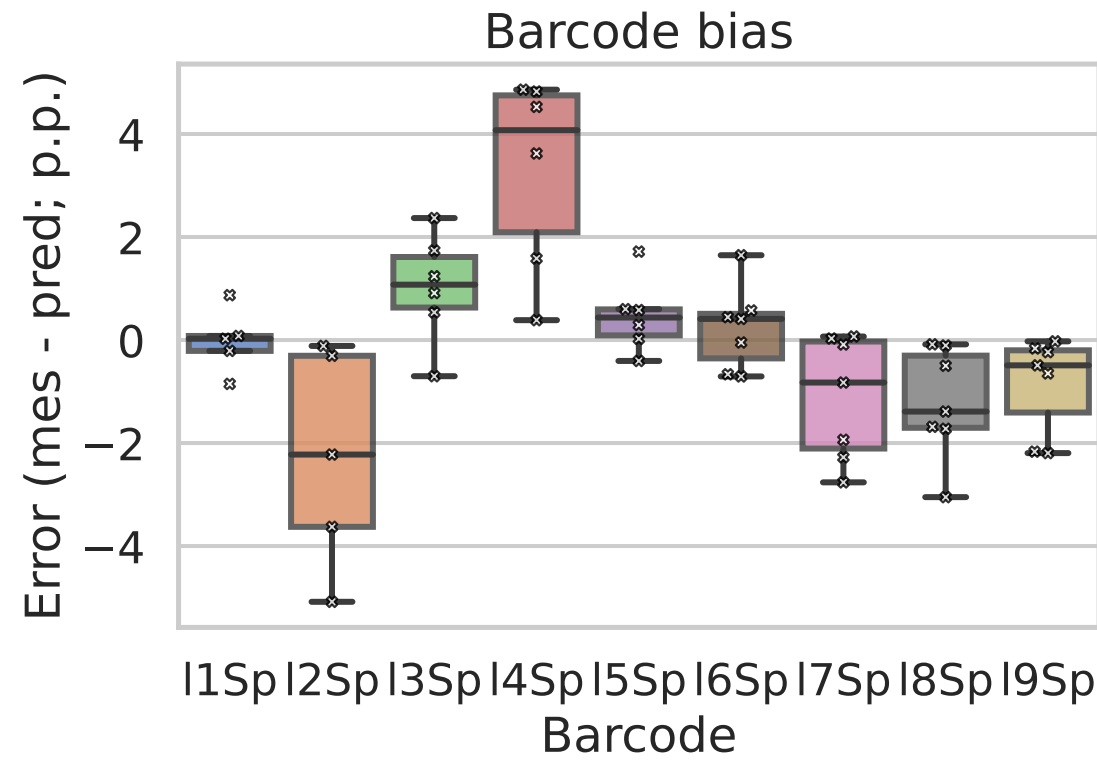
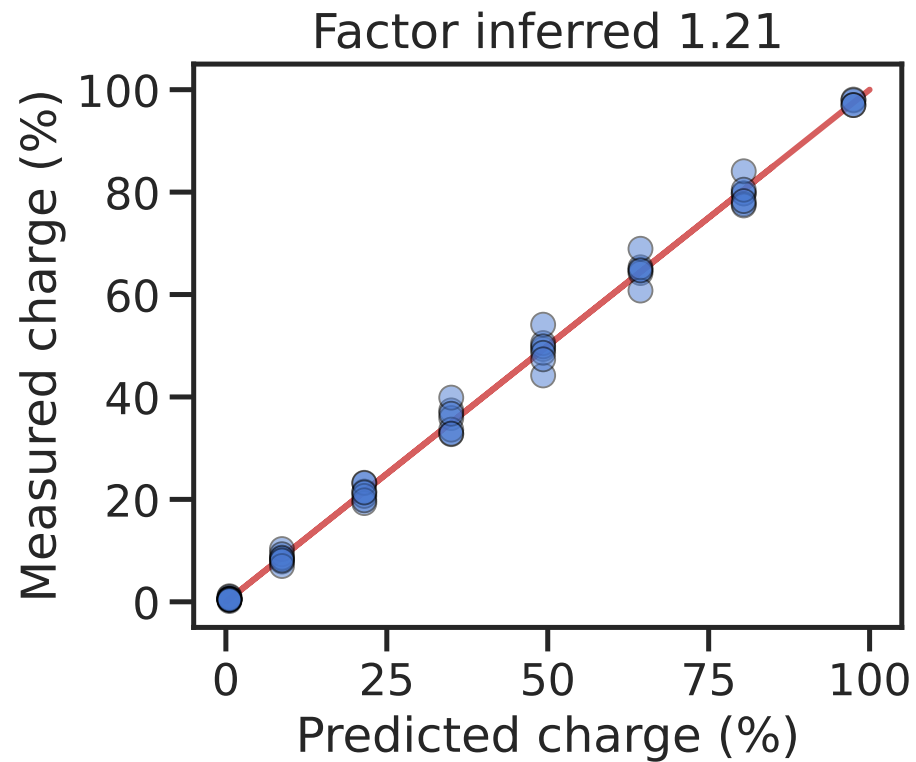
mito-Asp-GTC



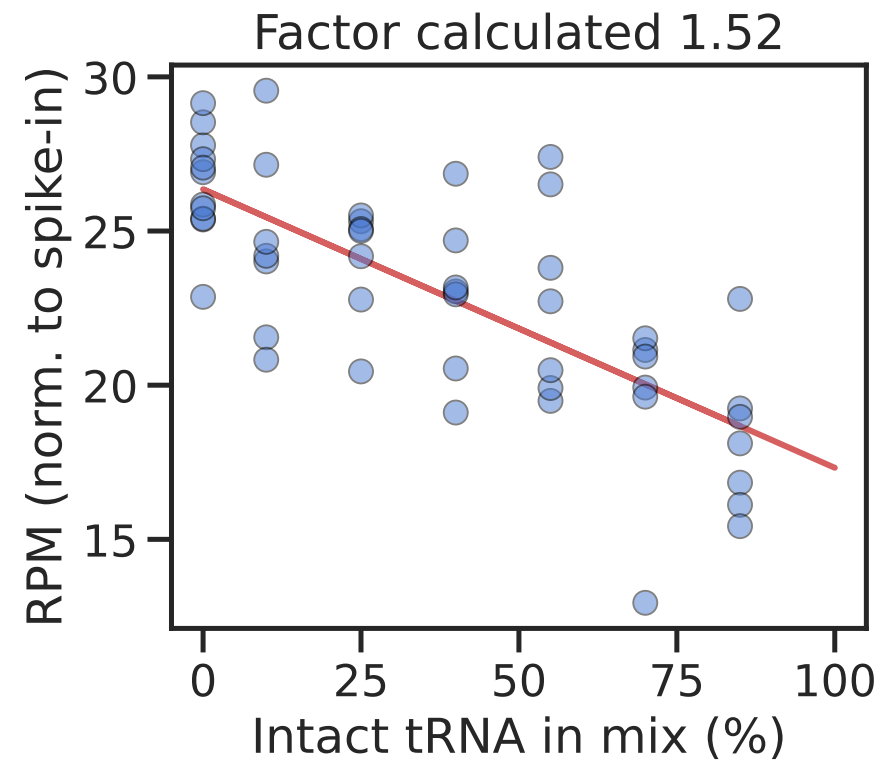
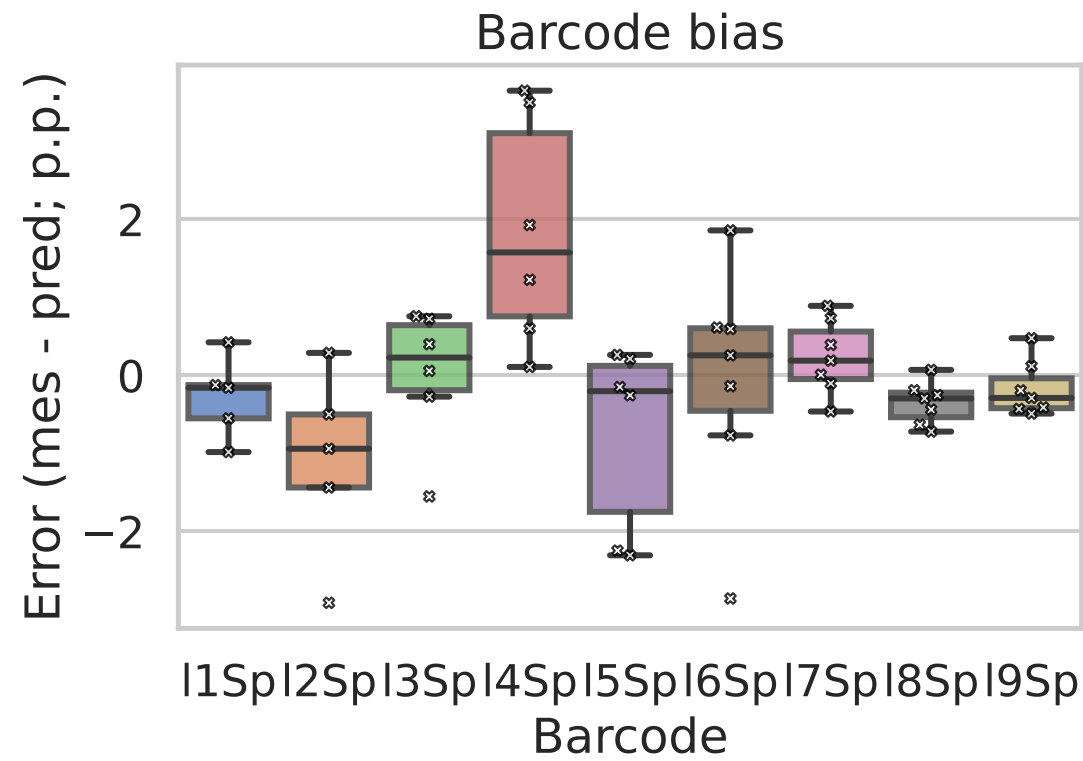
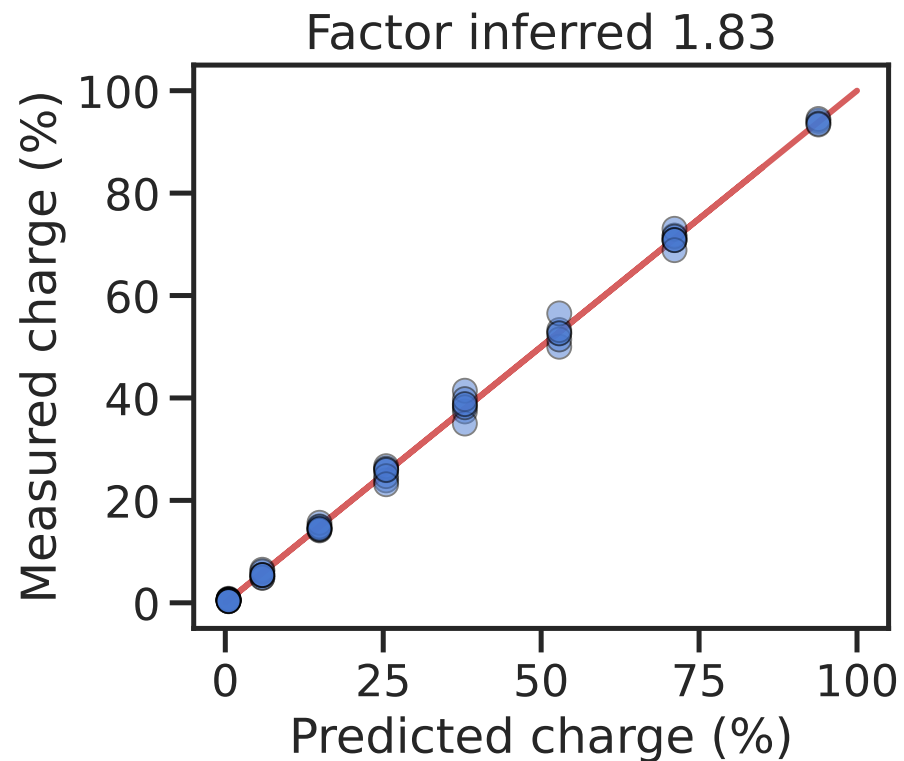
Phe-GAA-1-3



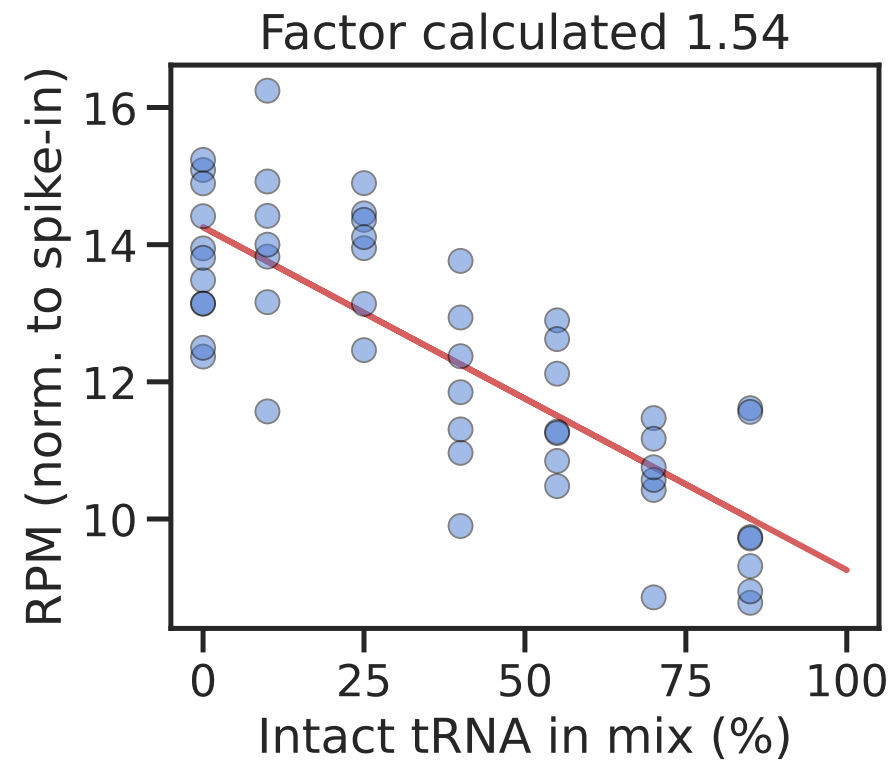
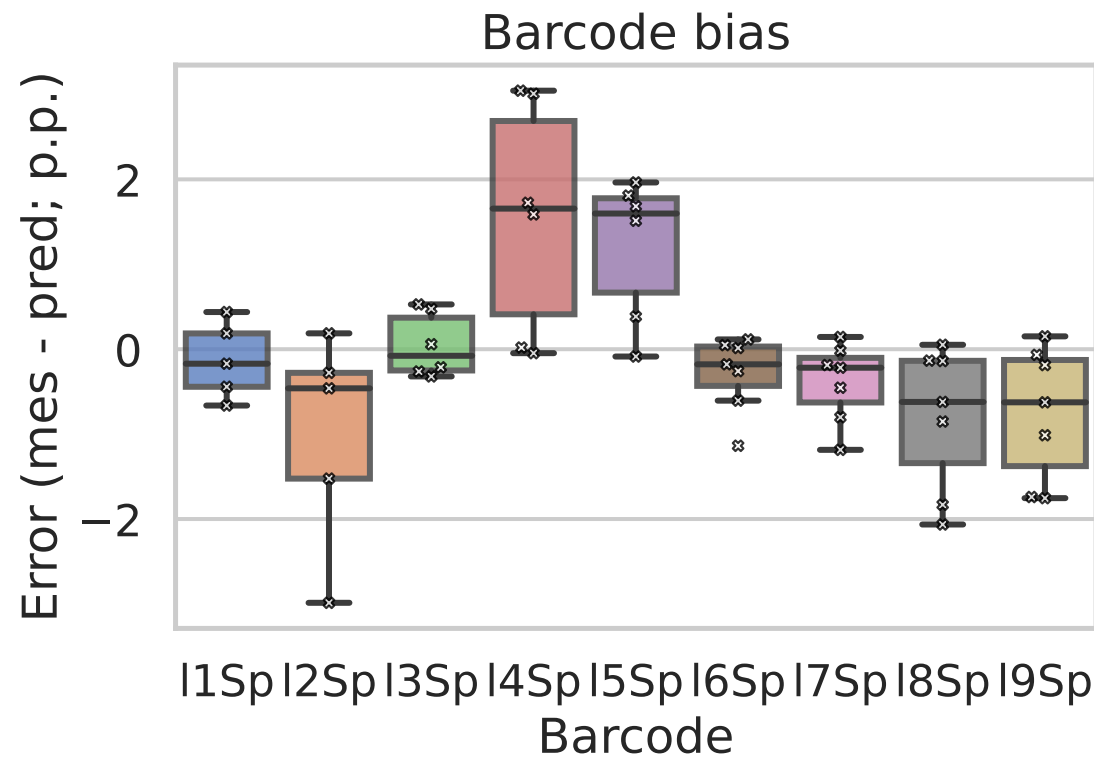
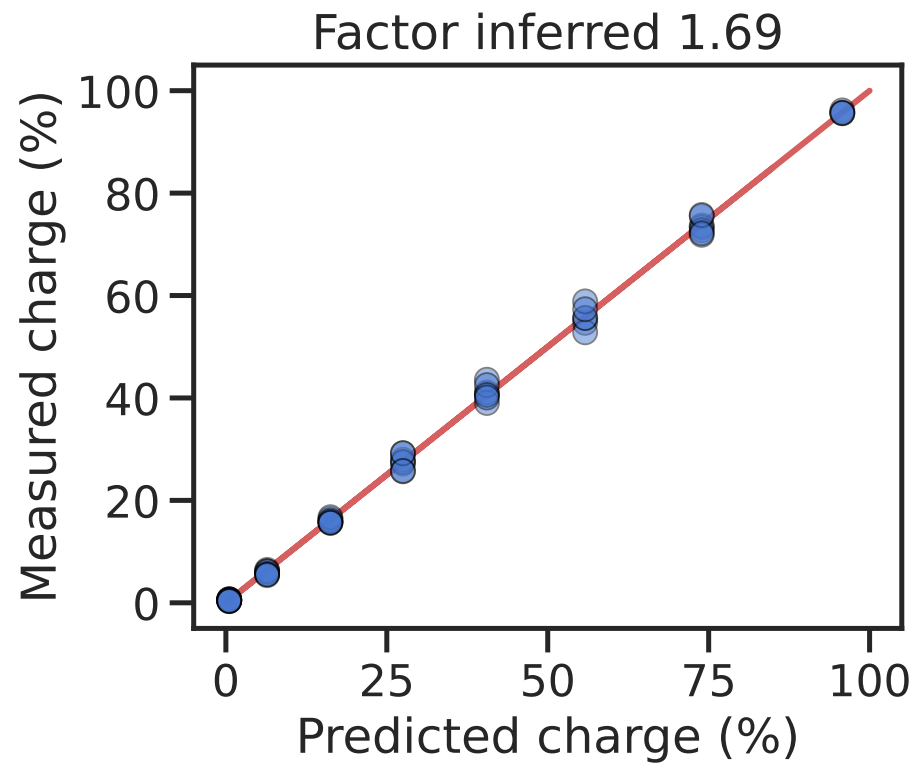
Gln-TTG-3-1



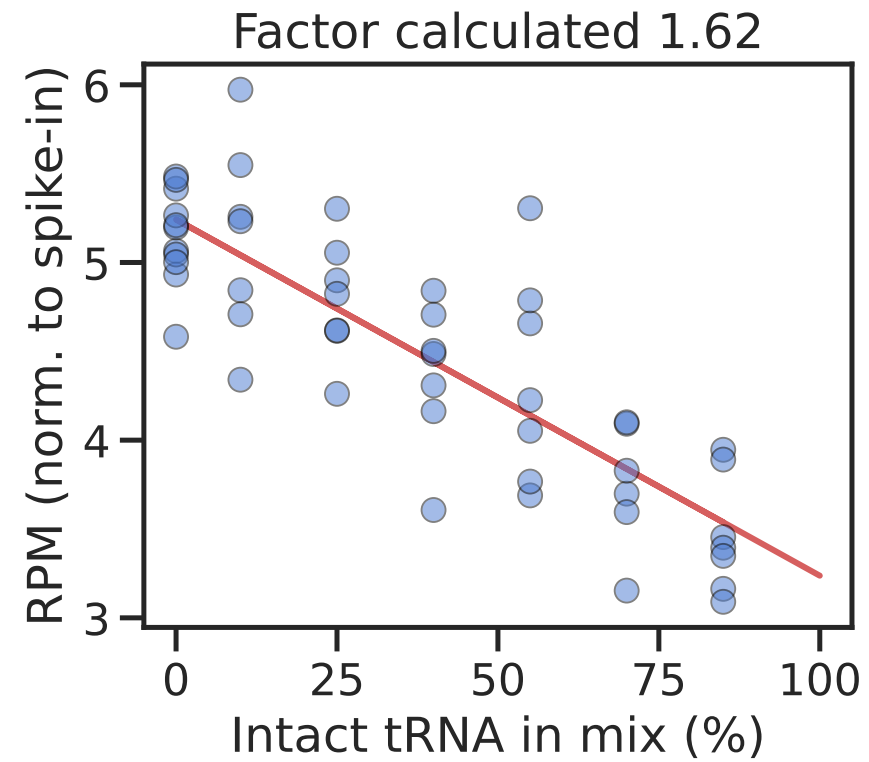
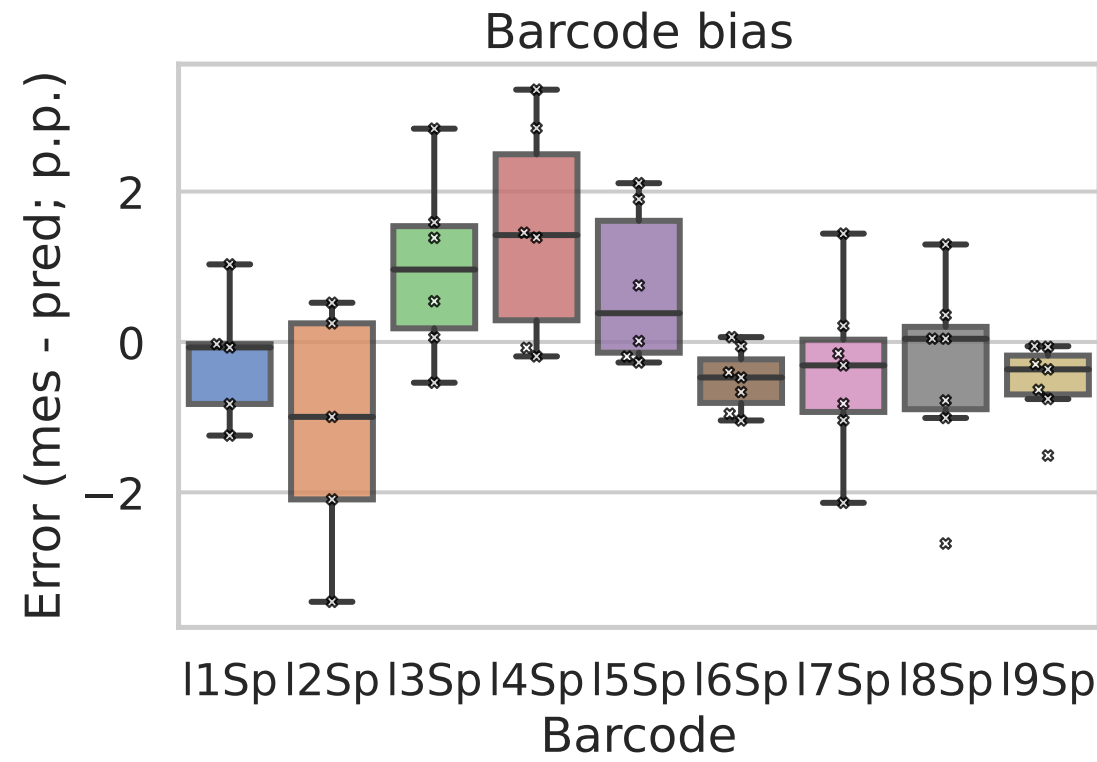
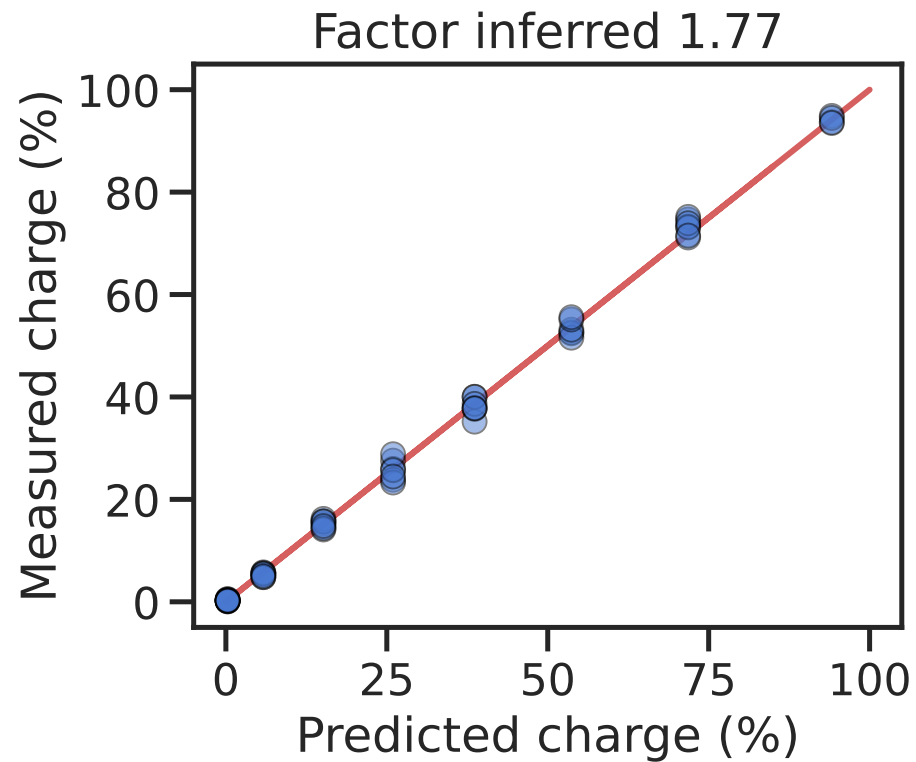
Leu-AAG-2-3



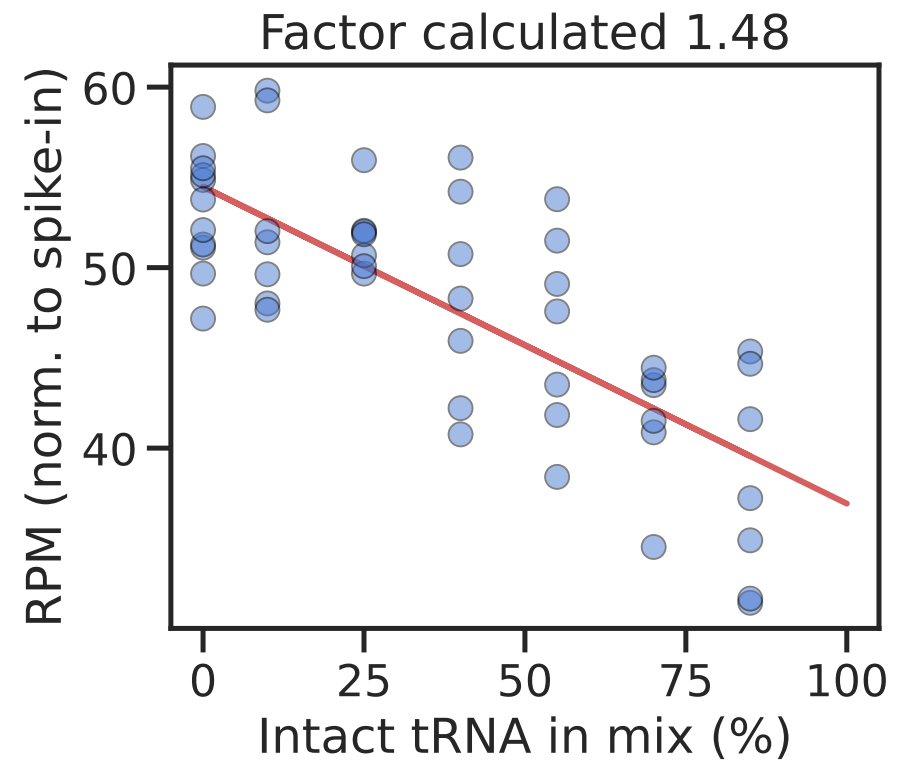
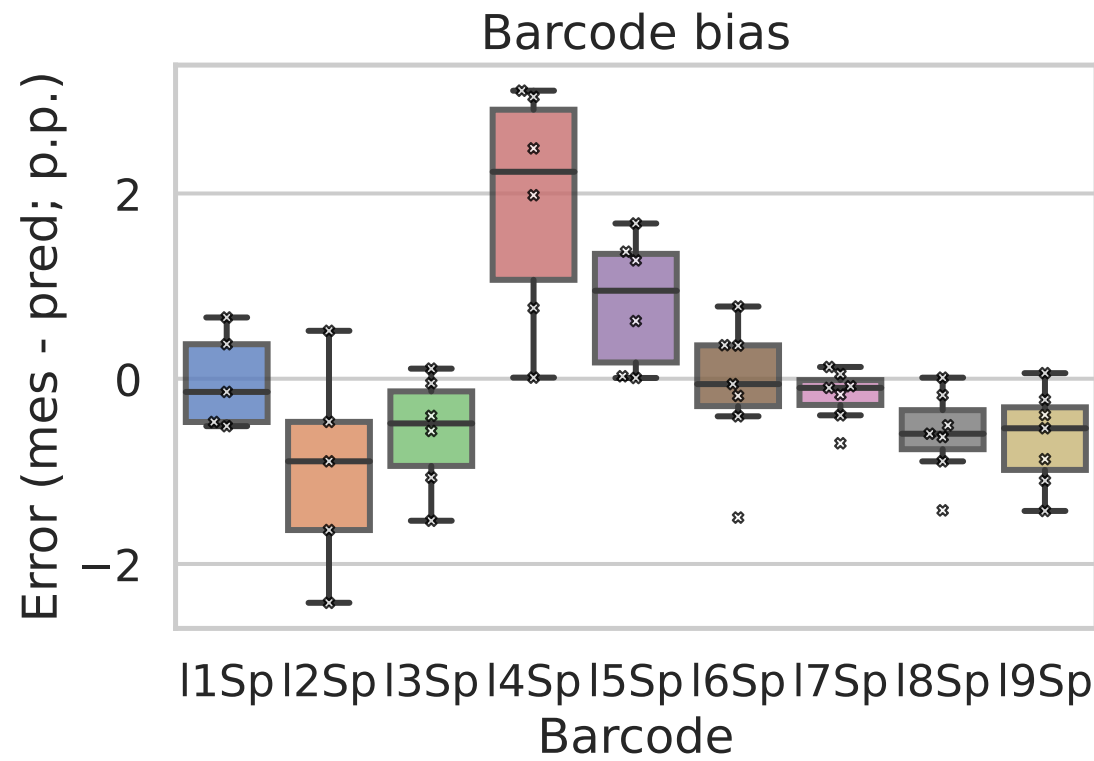
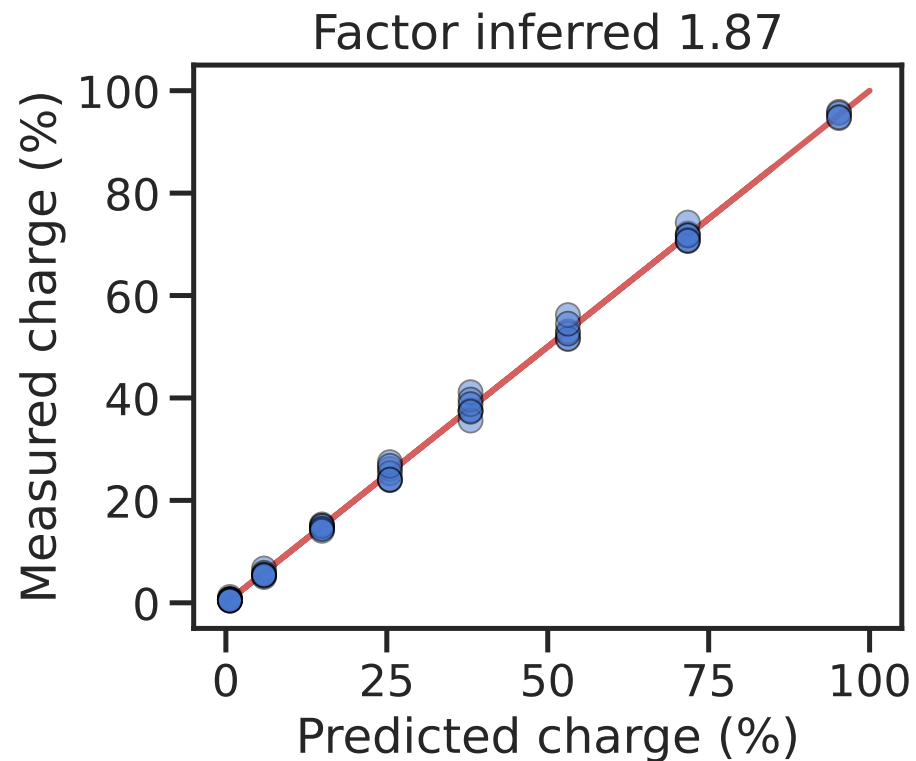
Arg-TCG-3-1



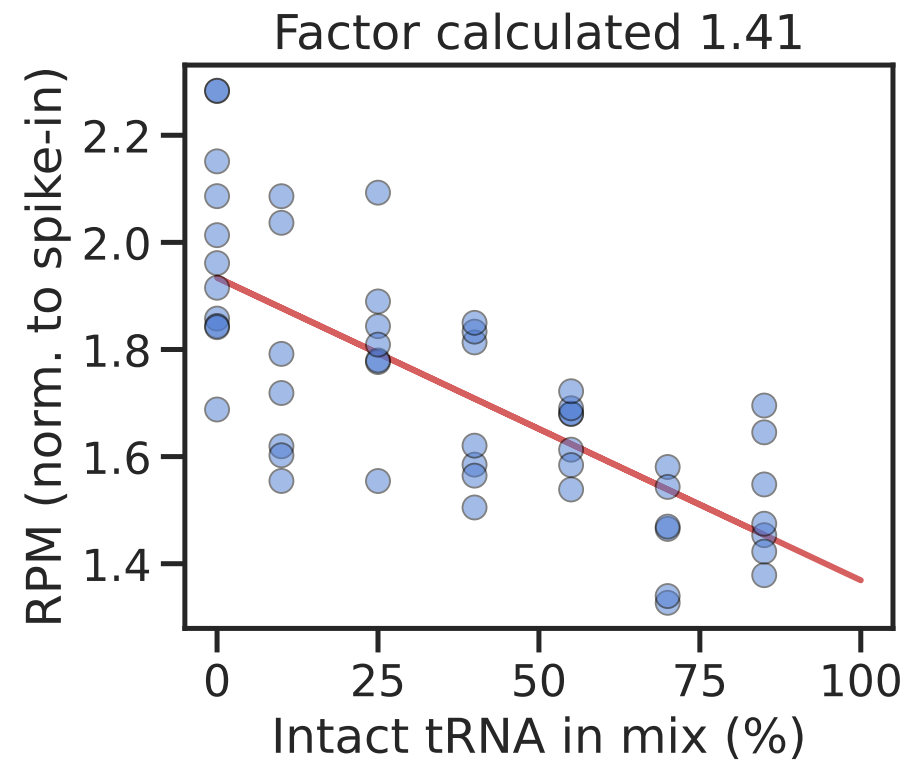
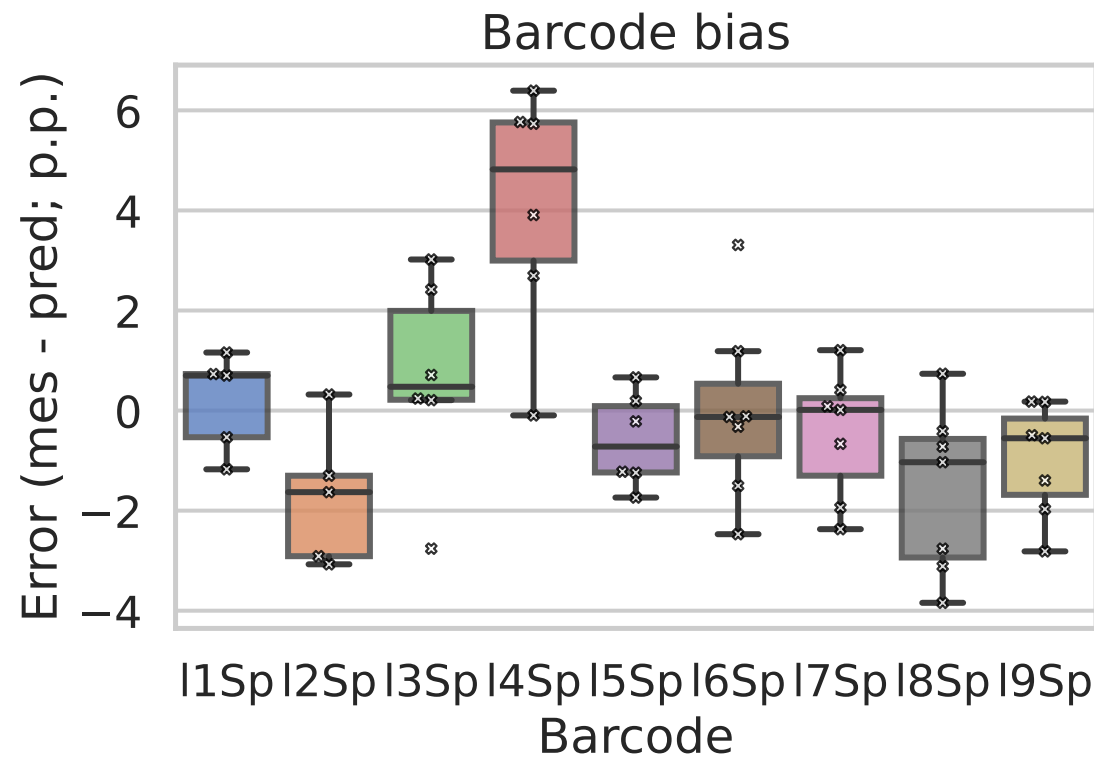
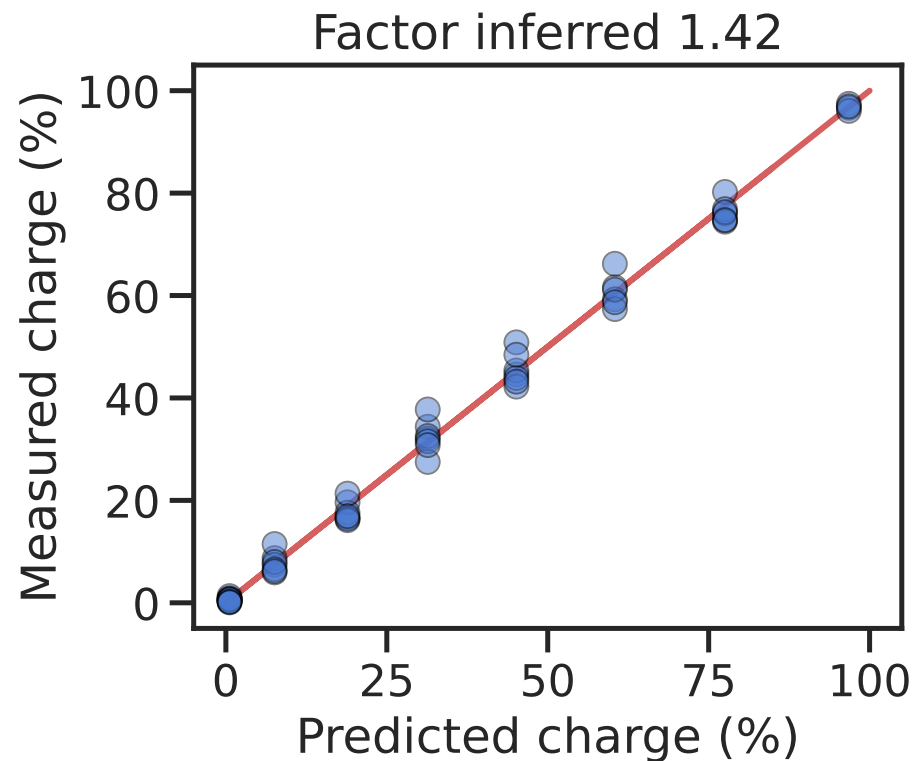
Pro-CGG-2-1



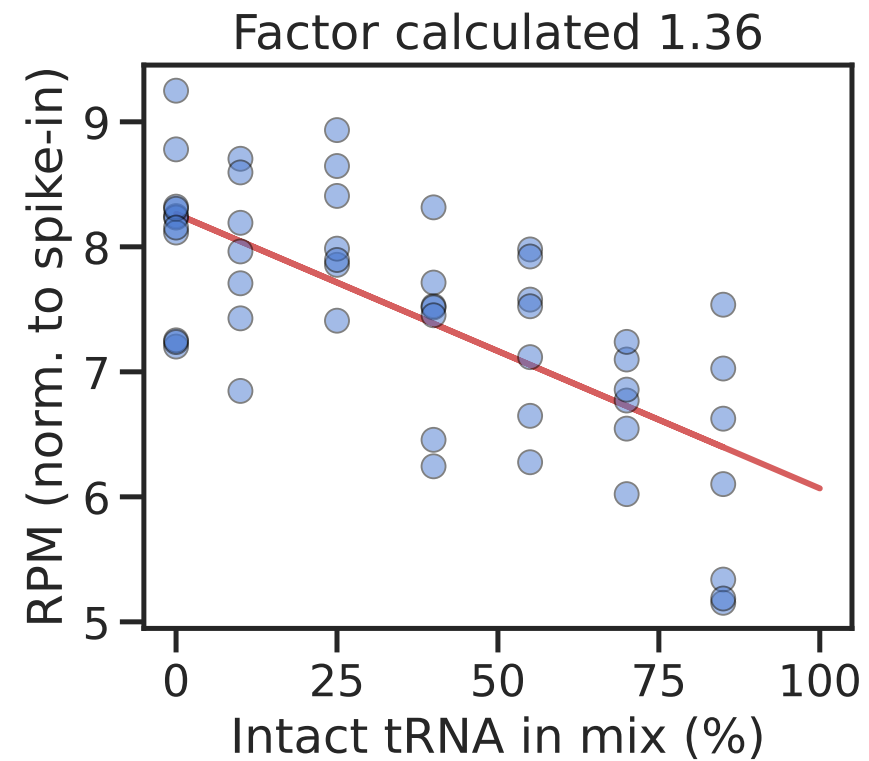
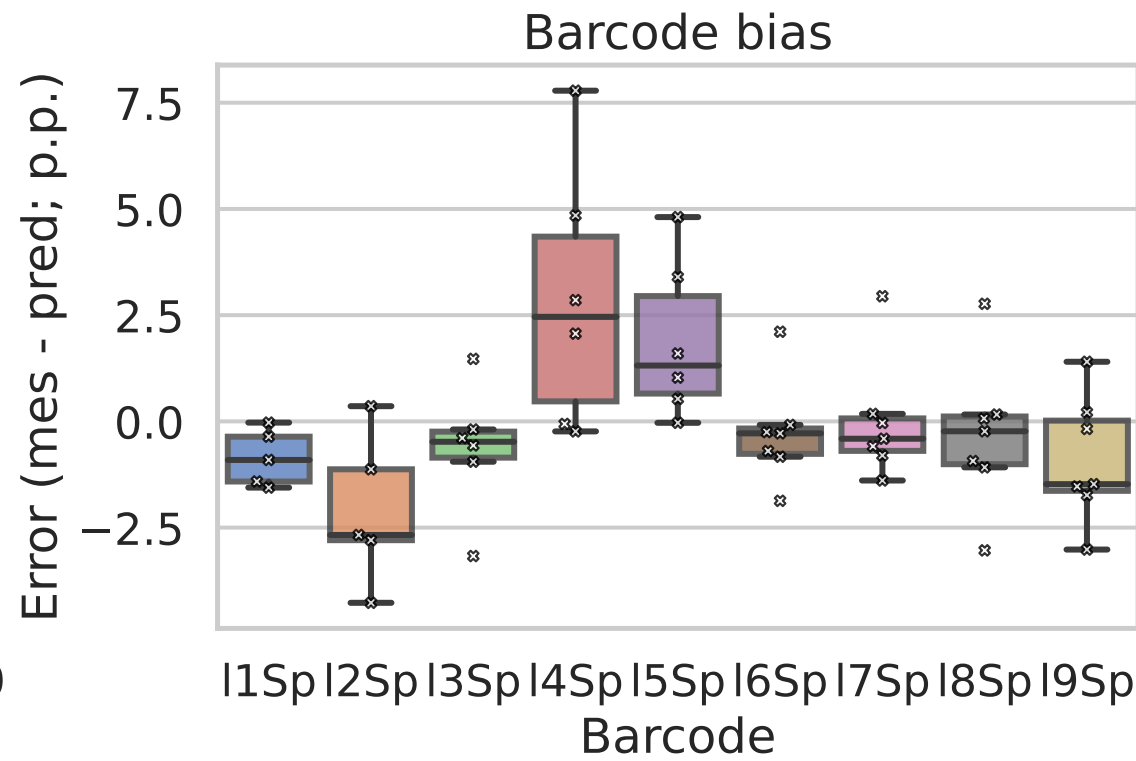
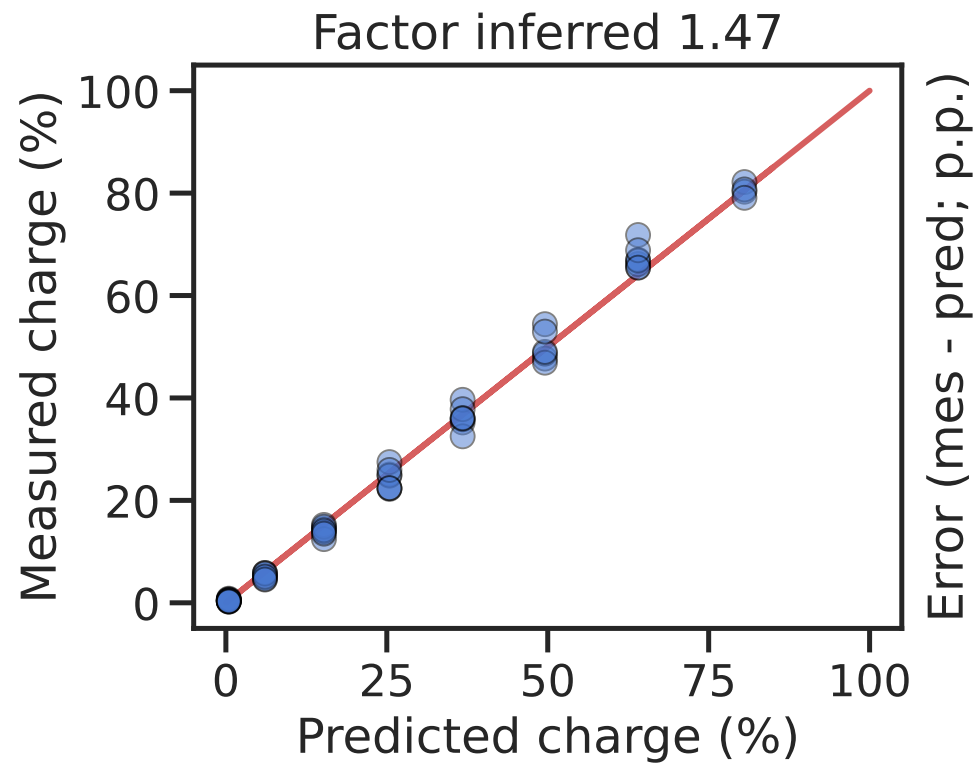
Leu-CAG-1-1



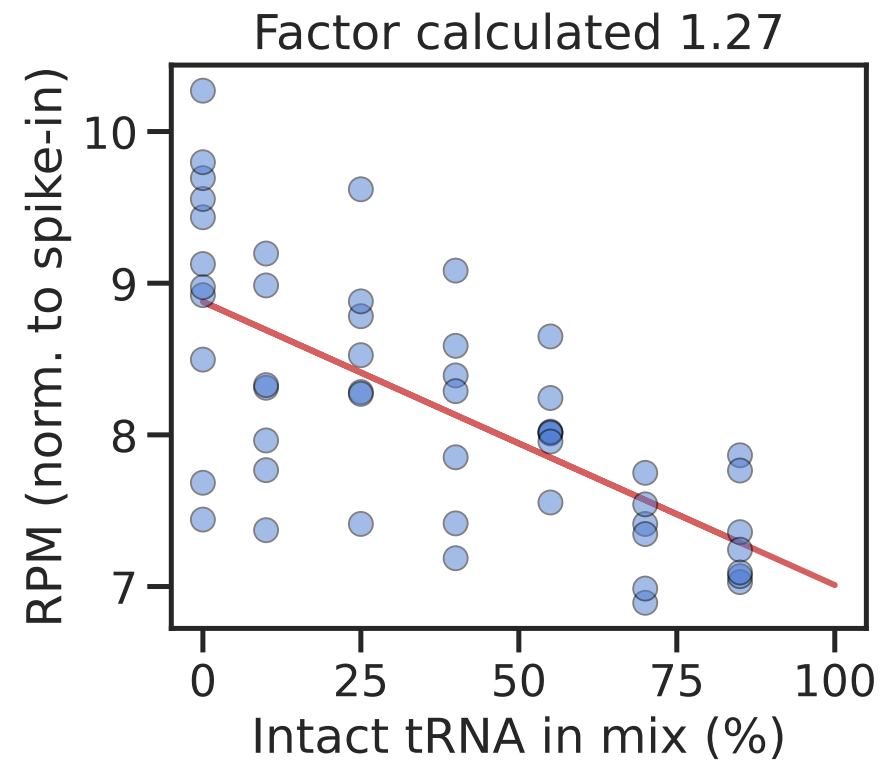
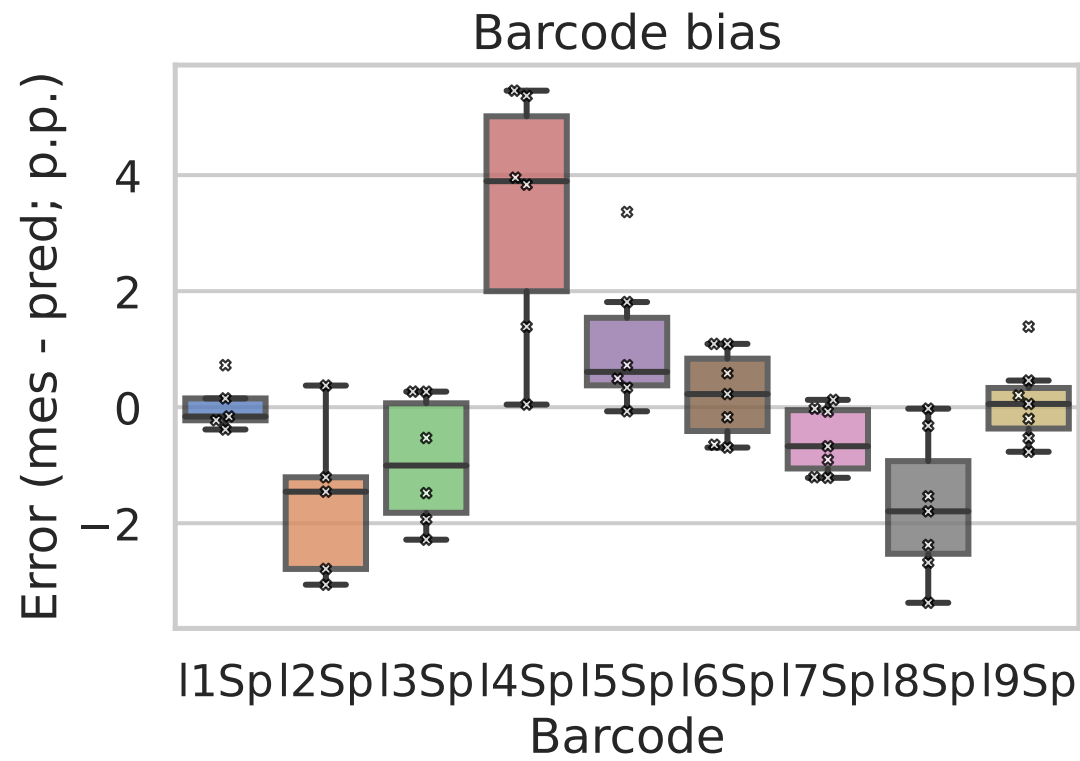
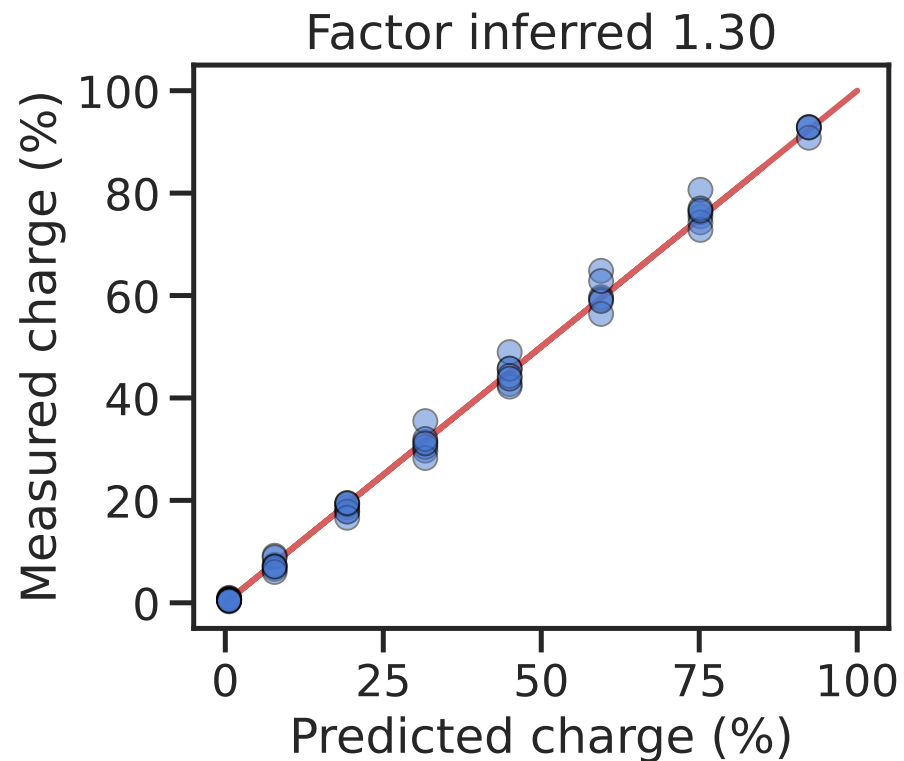
Gln-CTG-4-2



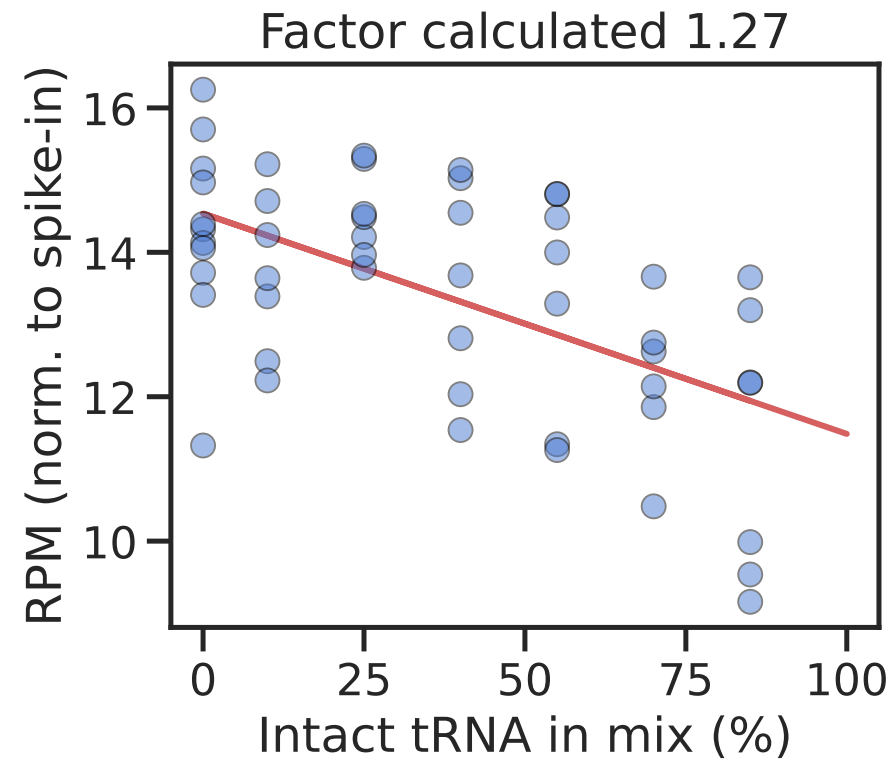
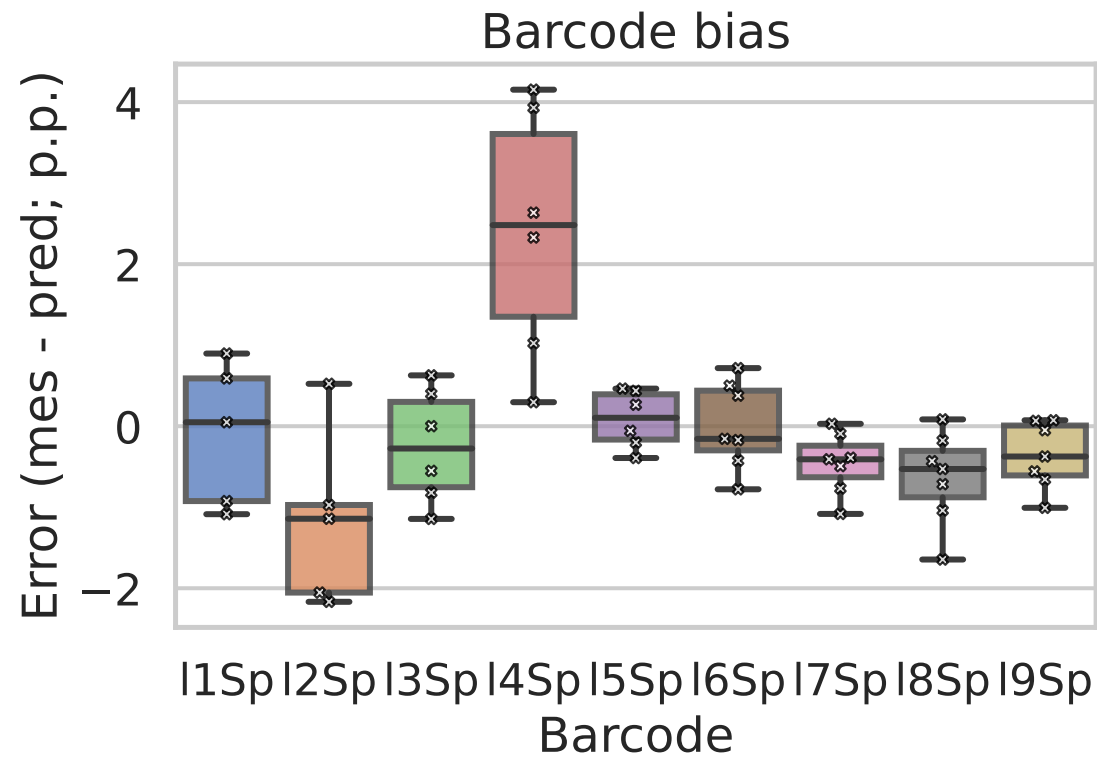
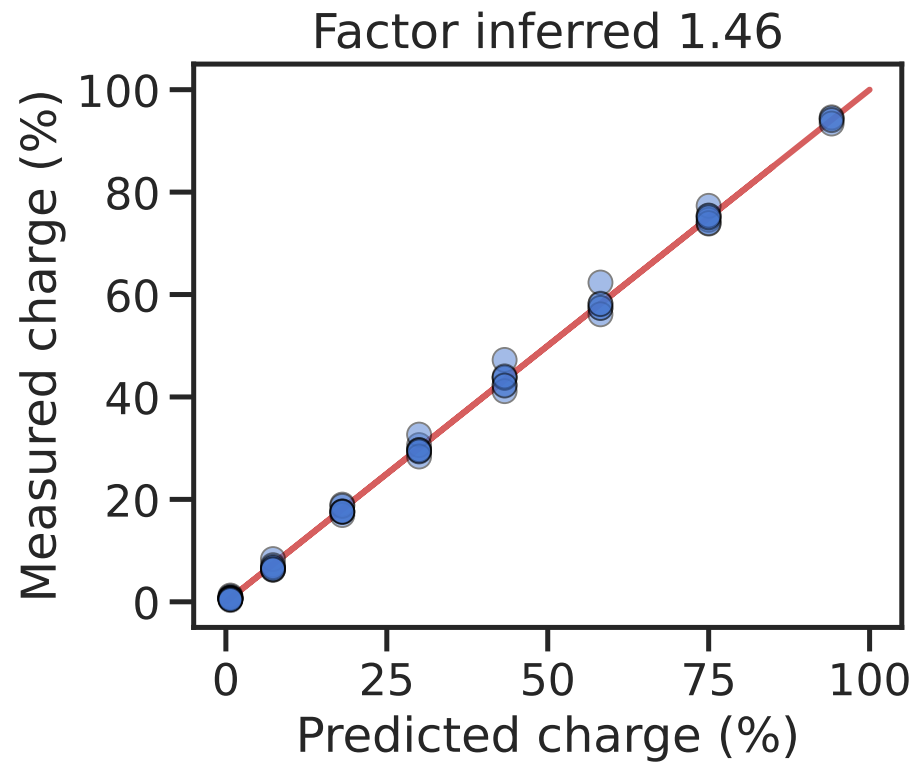
Ser-GCT-2-1



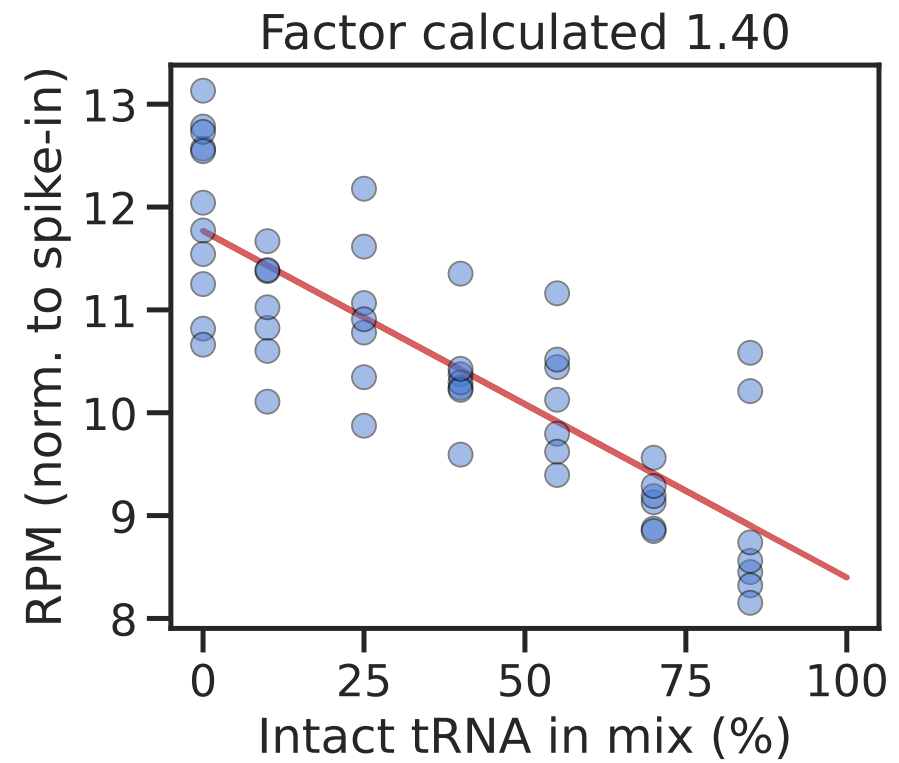
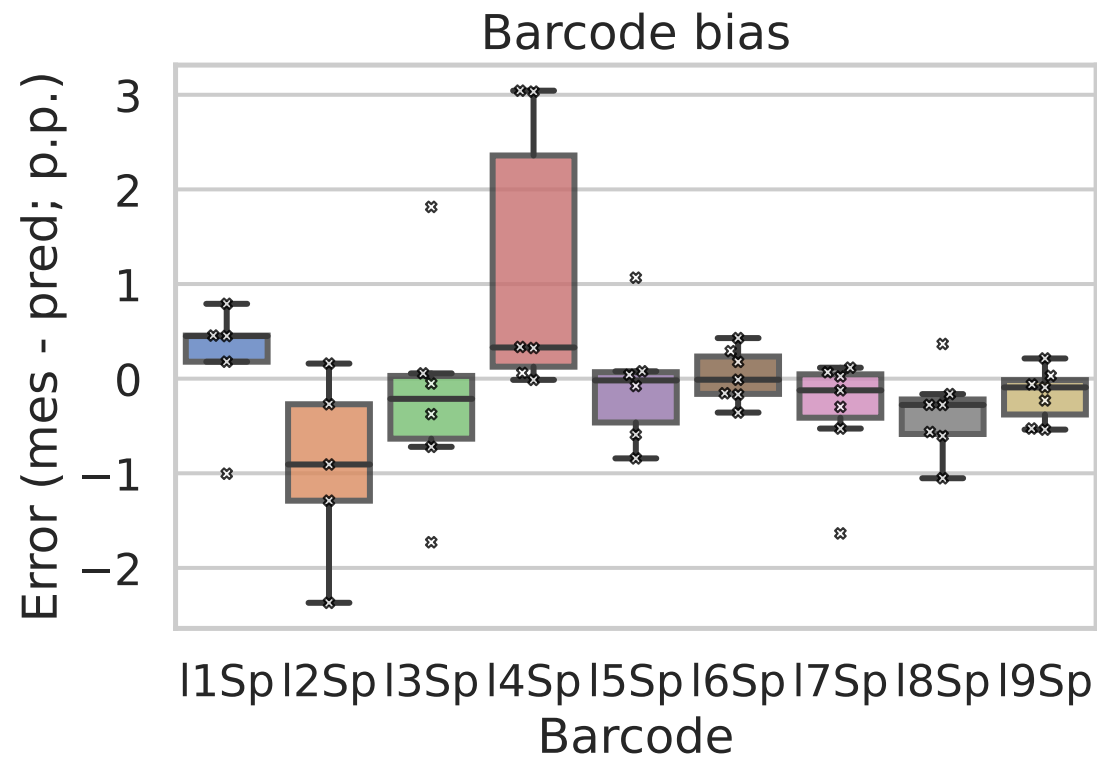
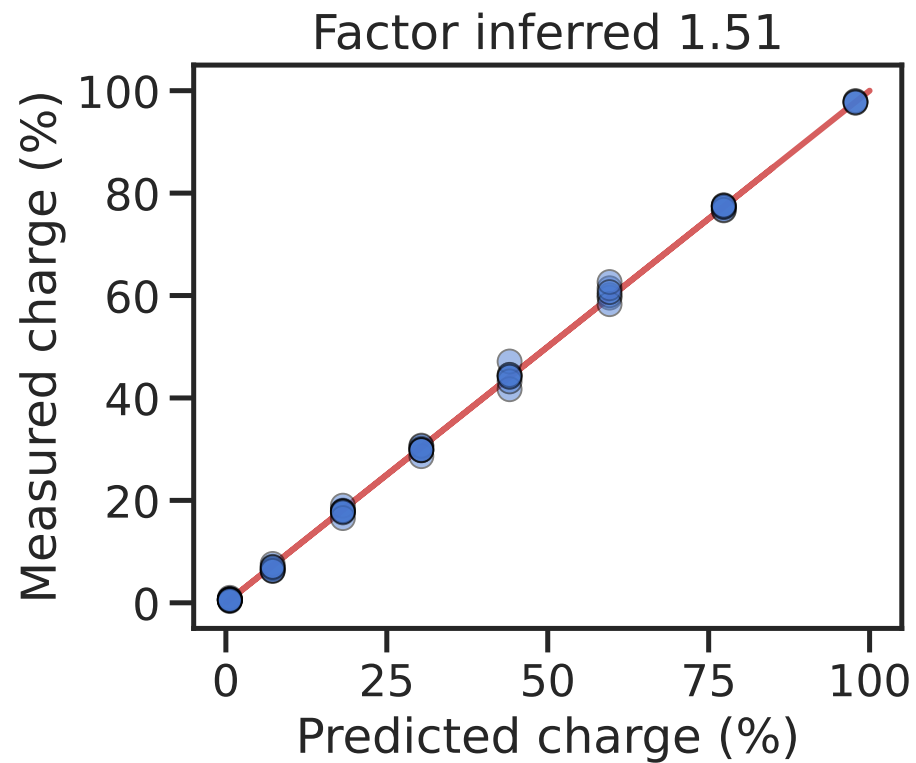
Thr-TGT-3-1



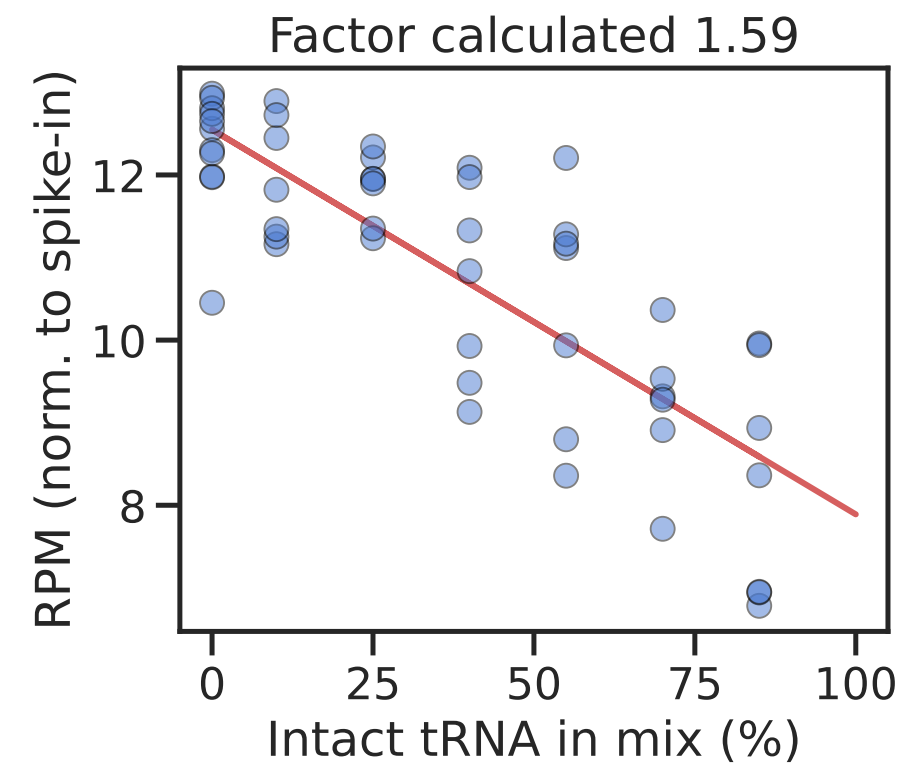
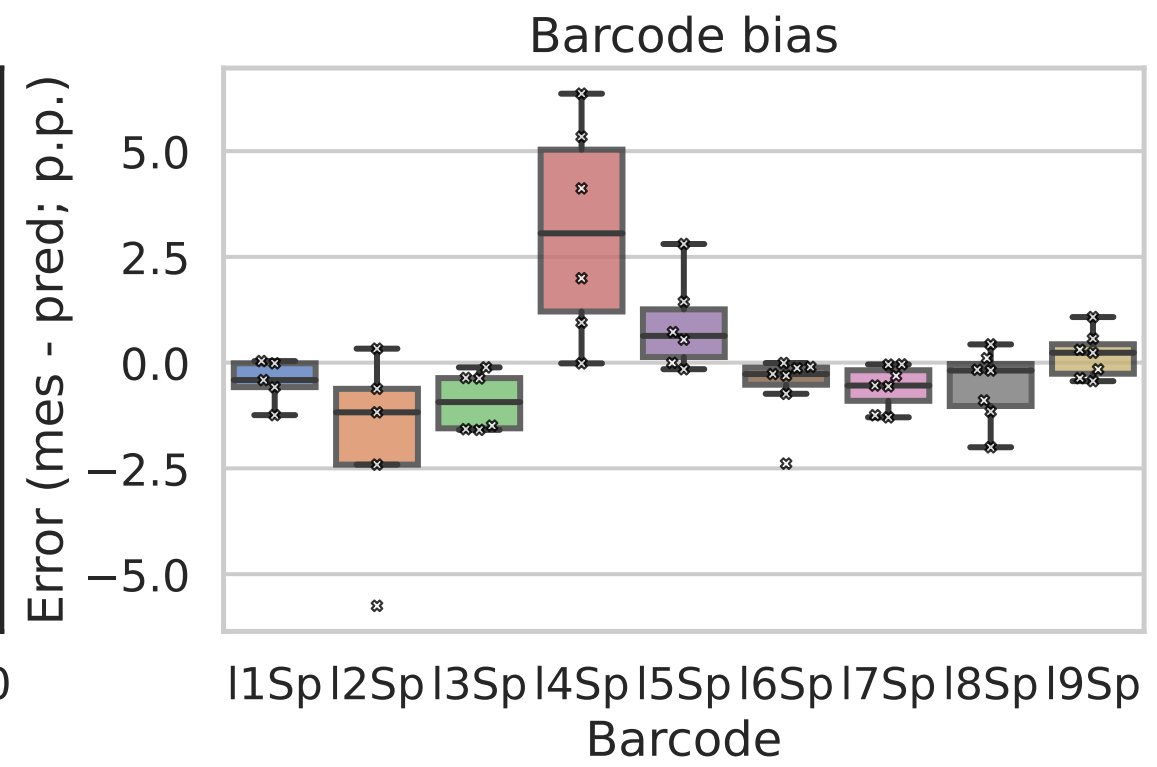
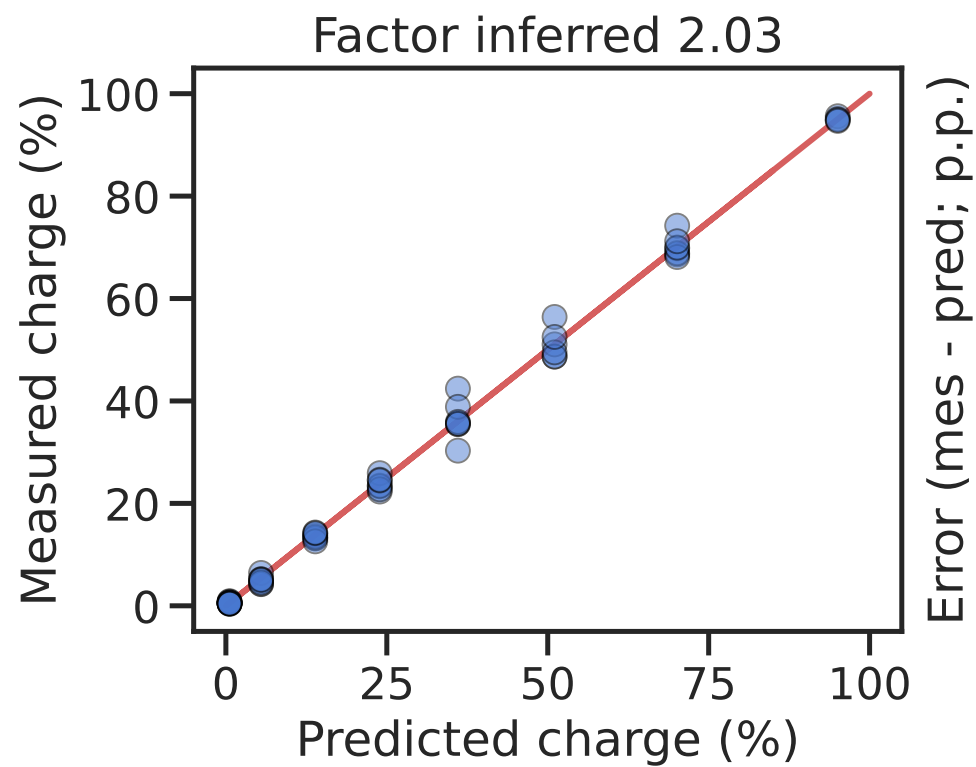
Ala-AGC-3-1



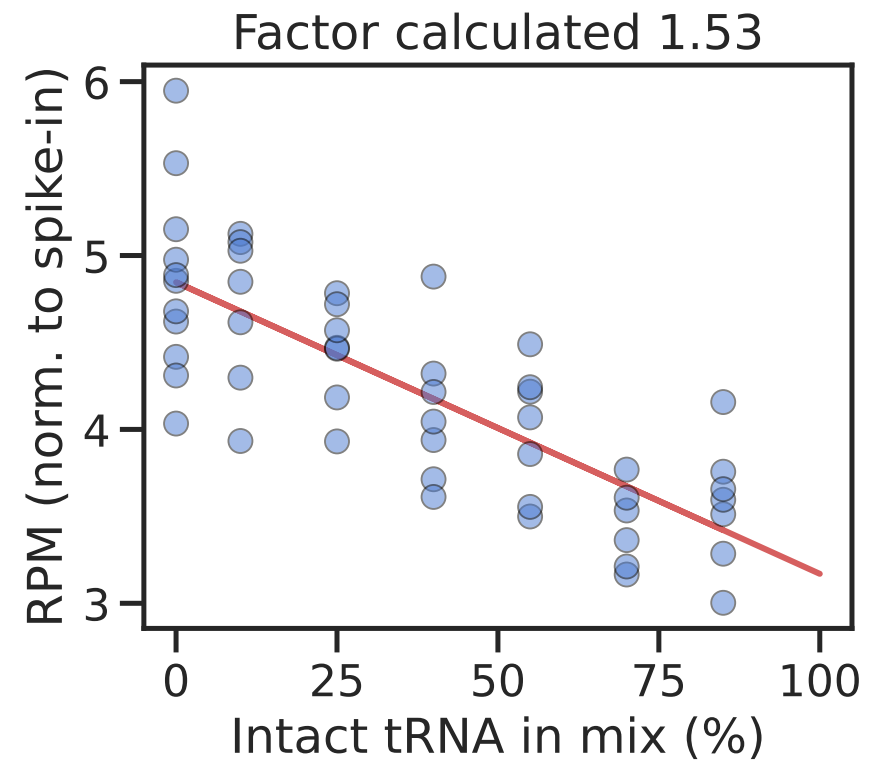
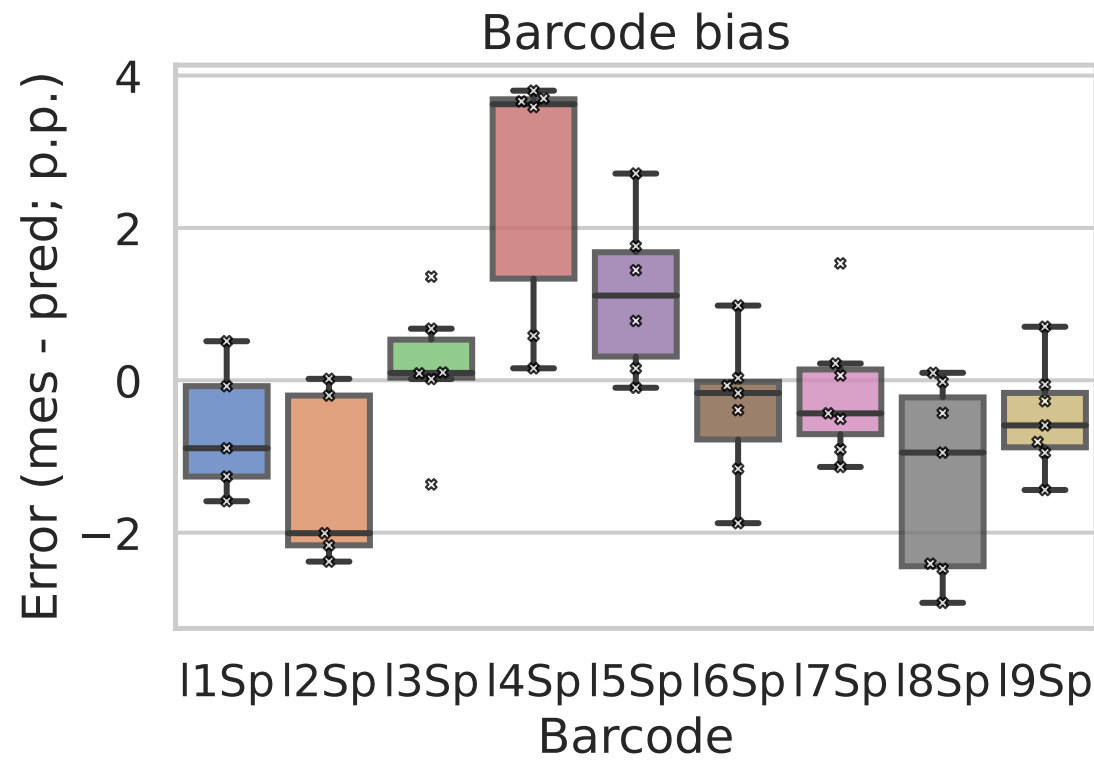
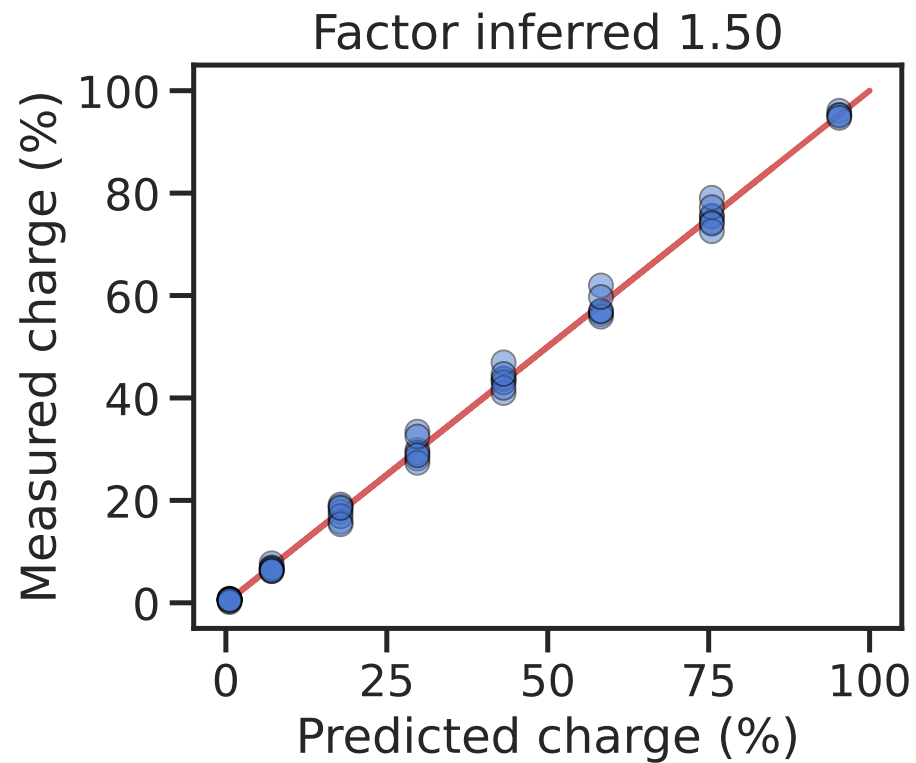
Ala-CGC-3-1



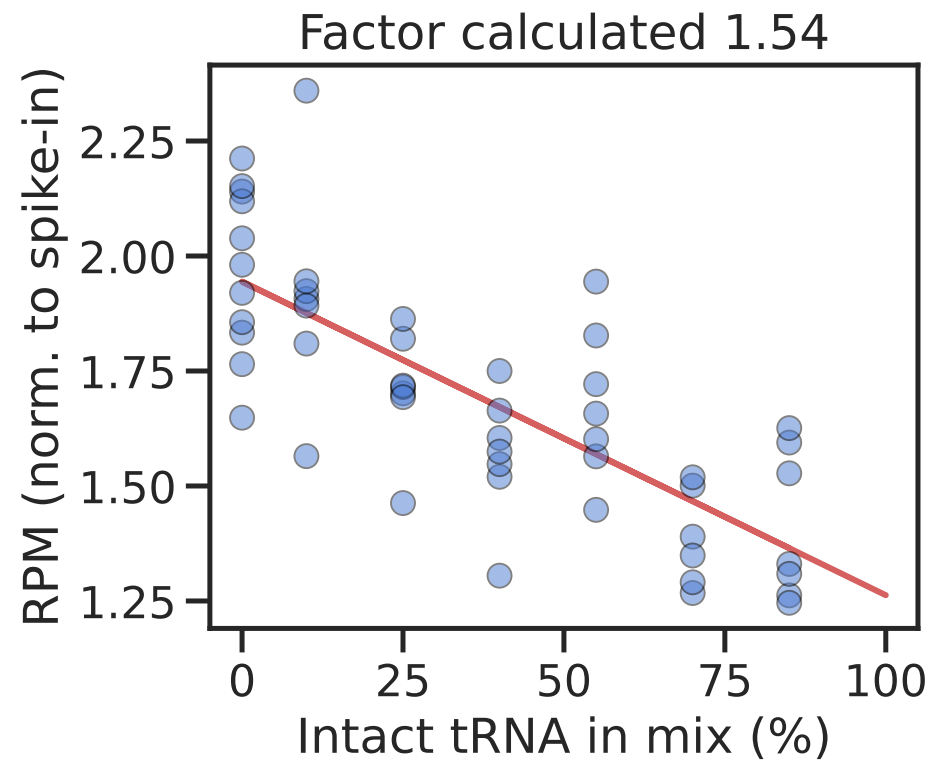
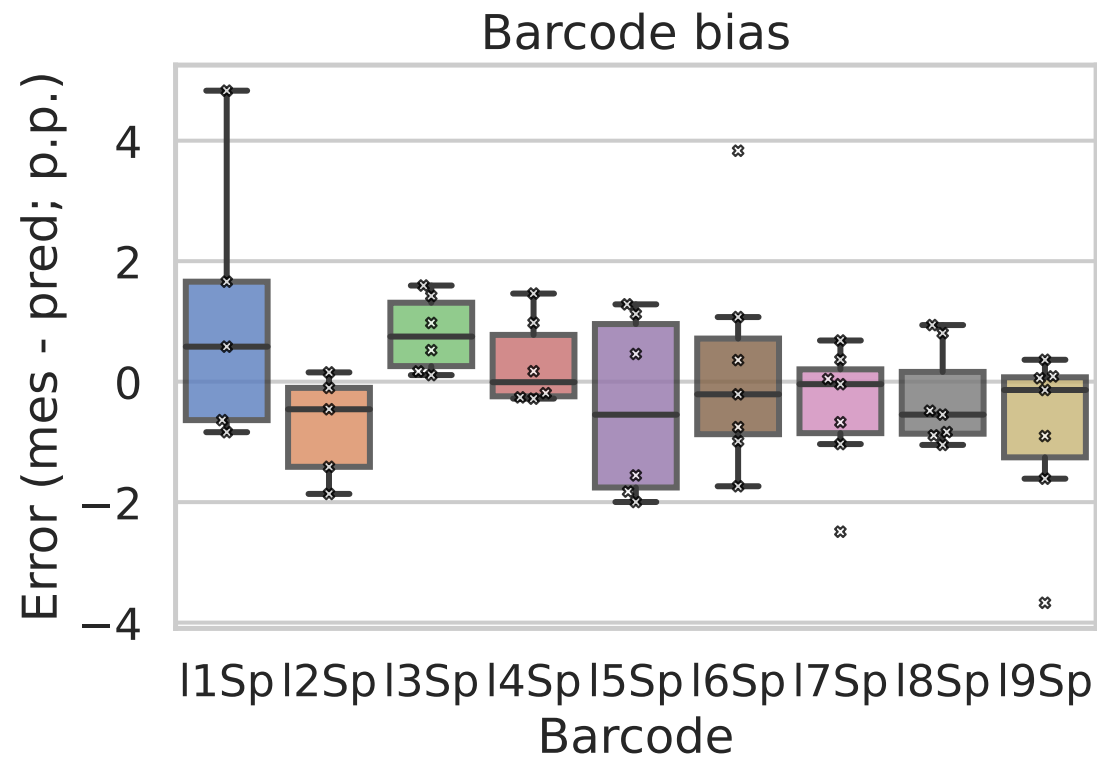
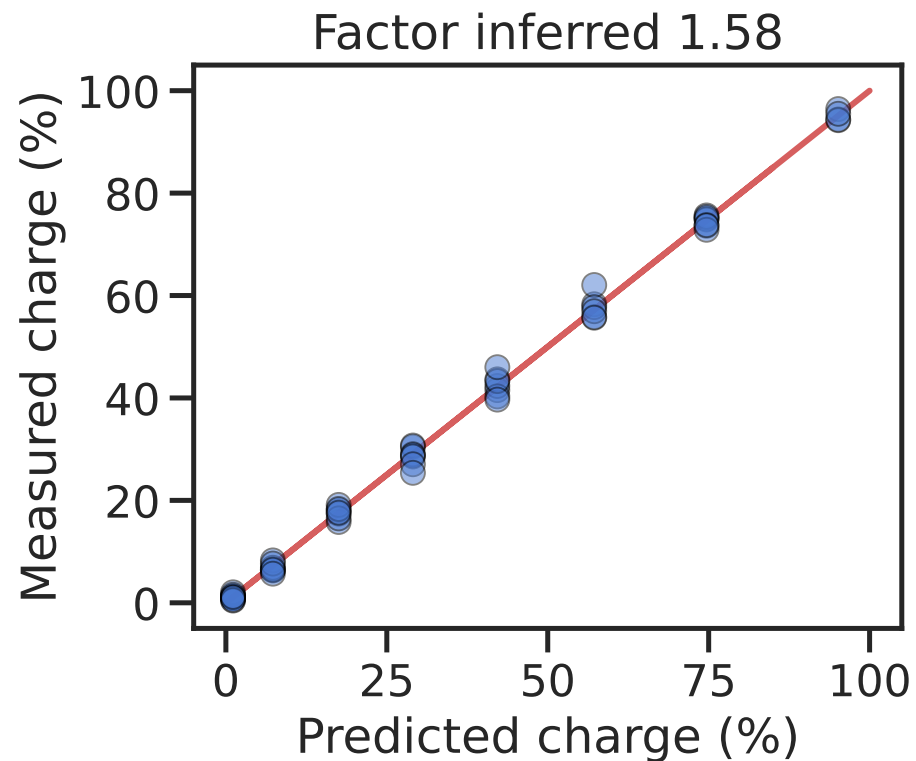
Leu-TAA-1-1



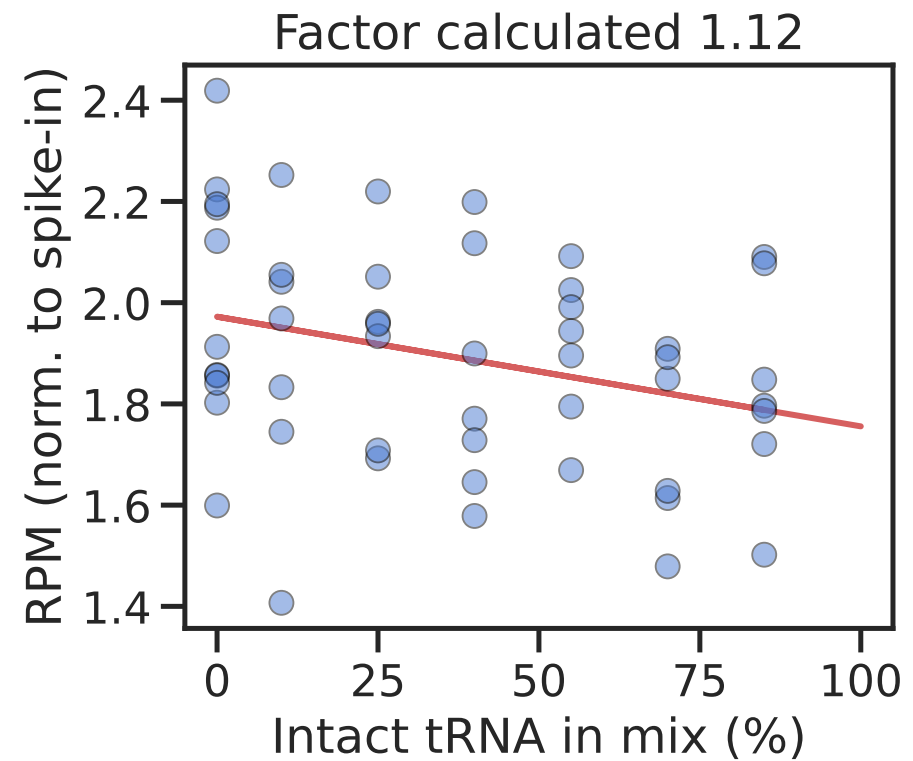
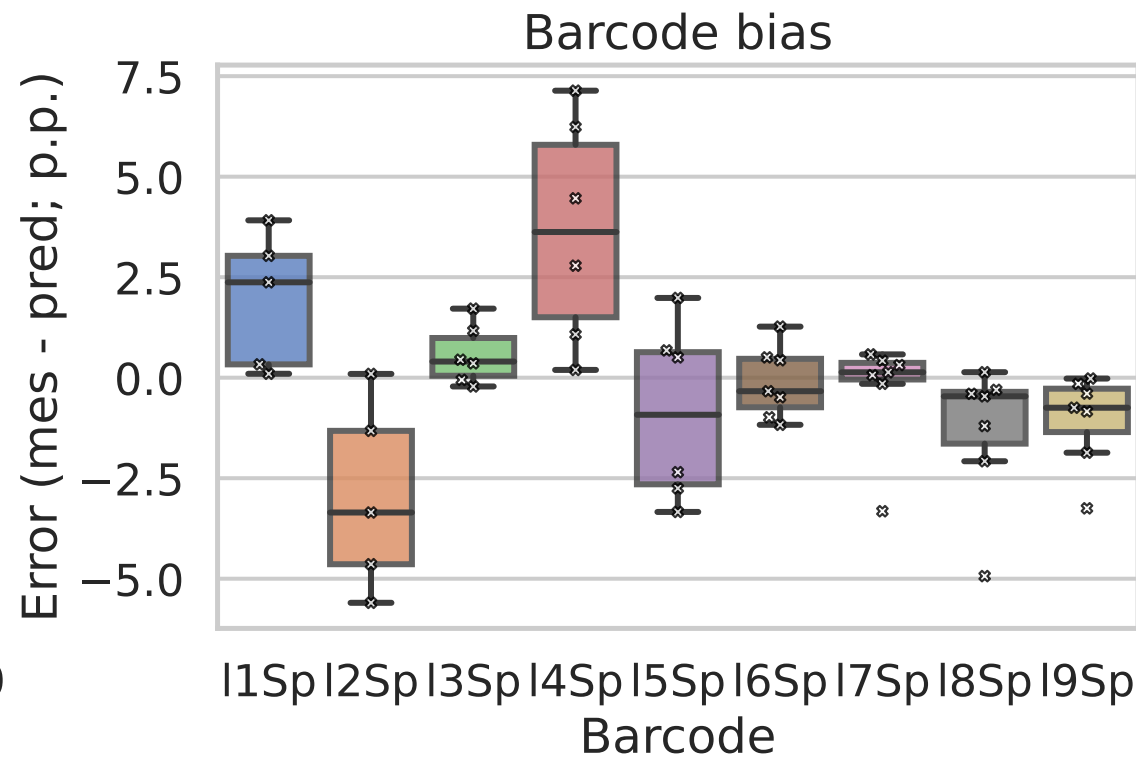
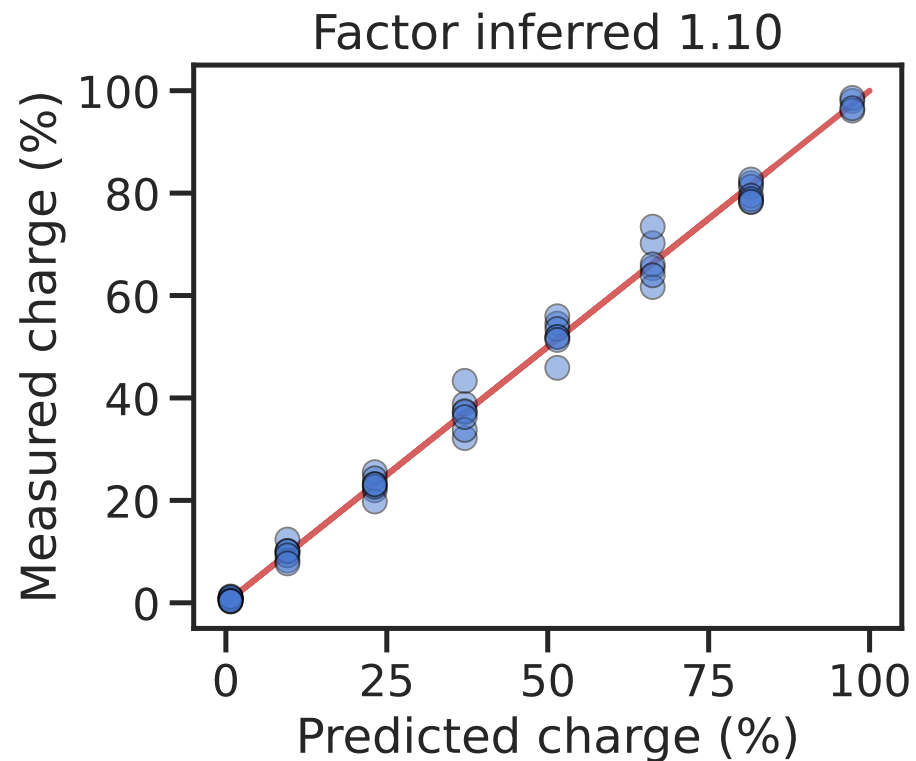
Lys-TTT-2-1



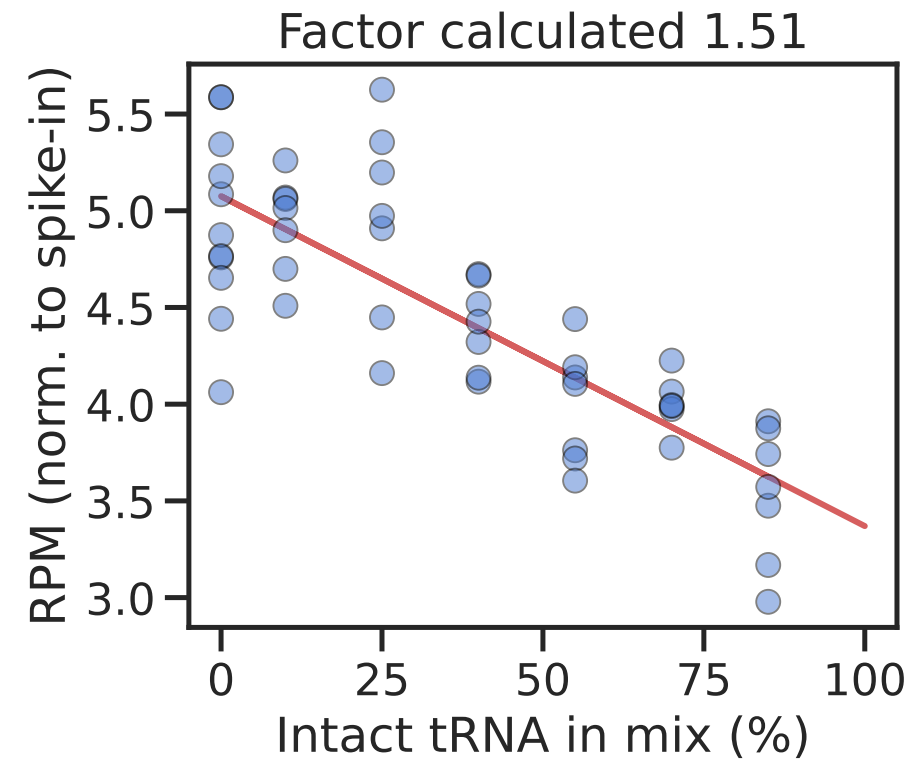
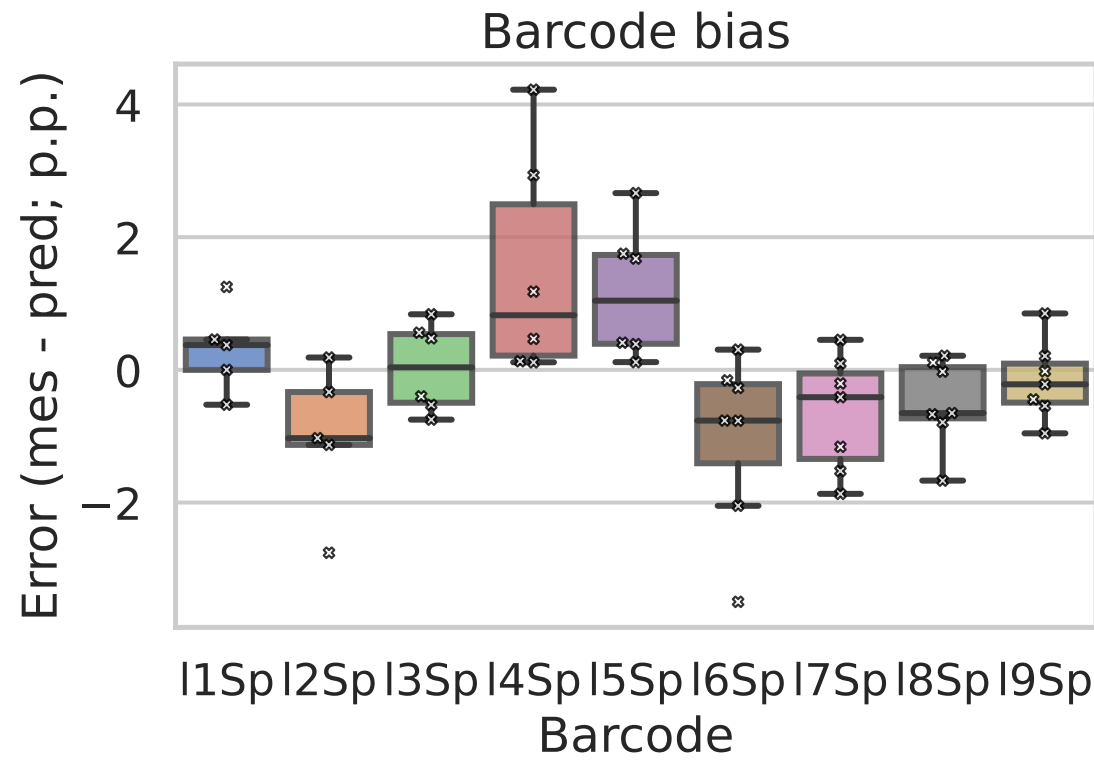
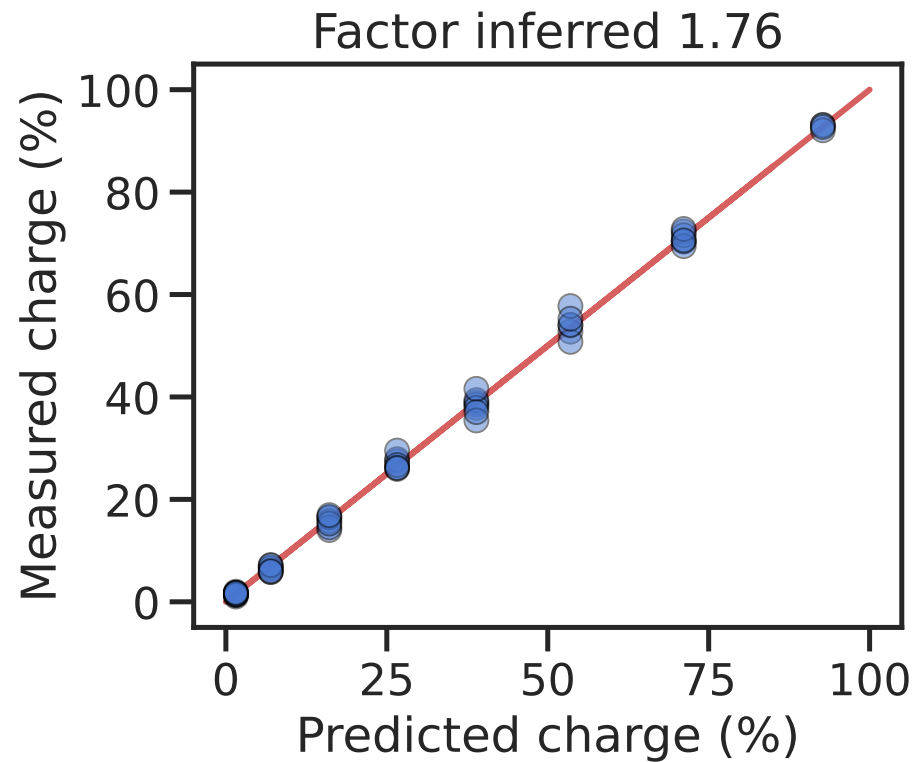
Ile-AAT-8-1



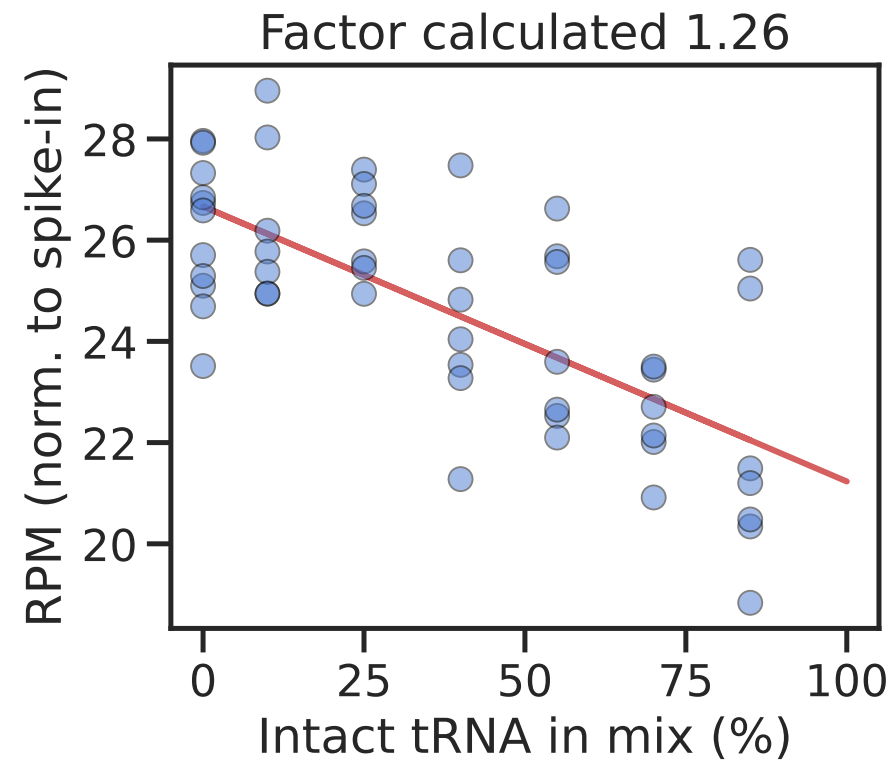
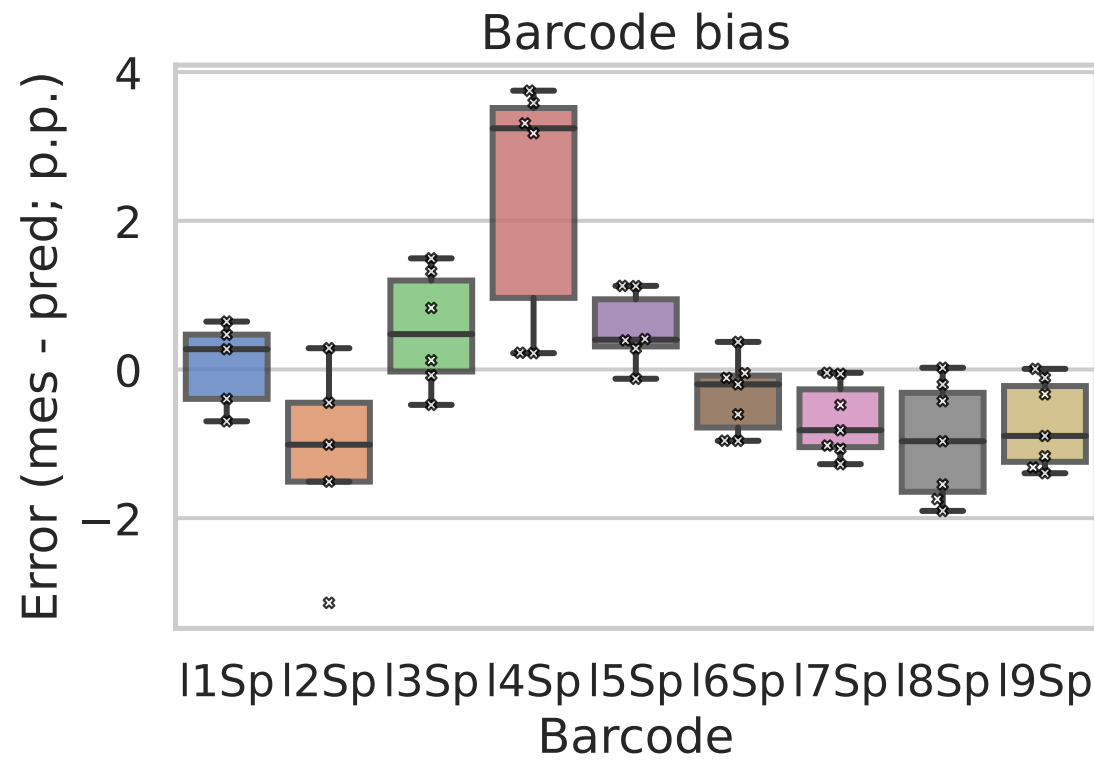
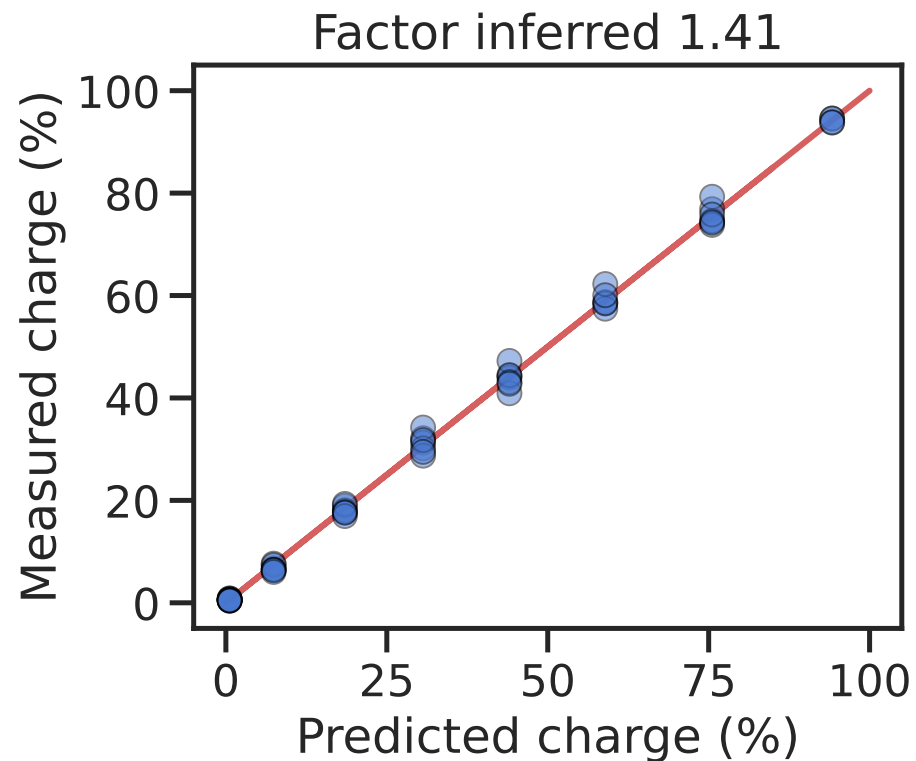
Cys-GCA-8-1



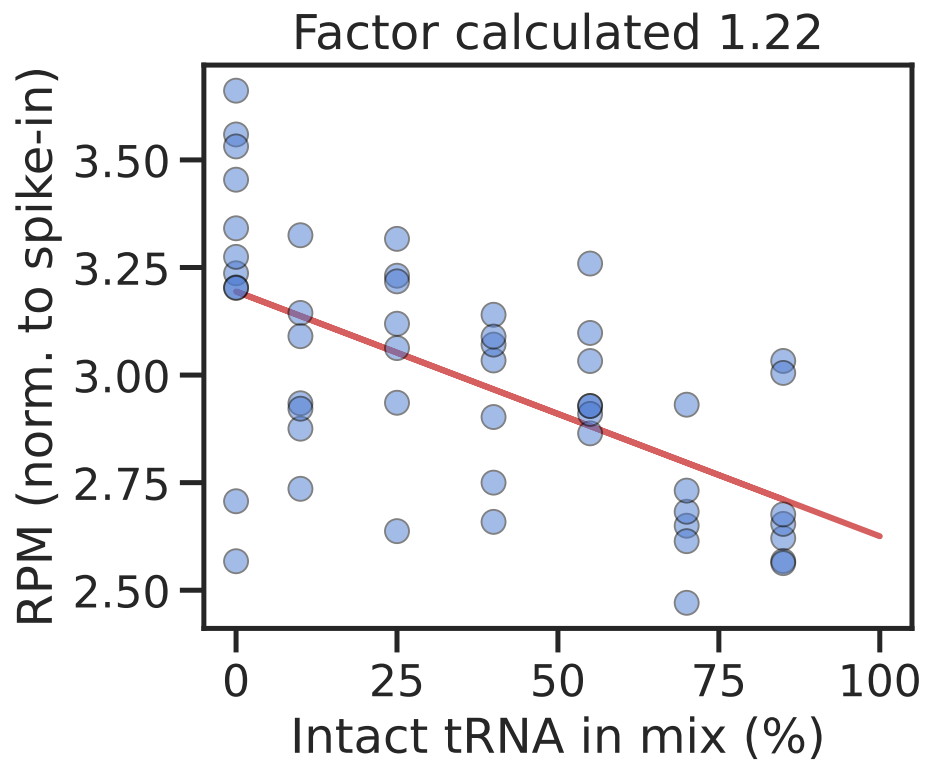
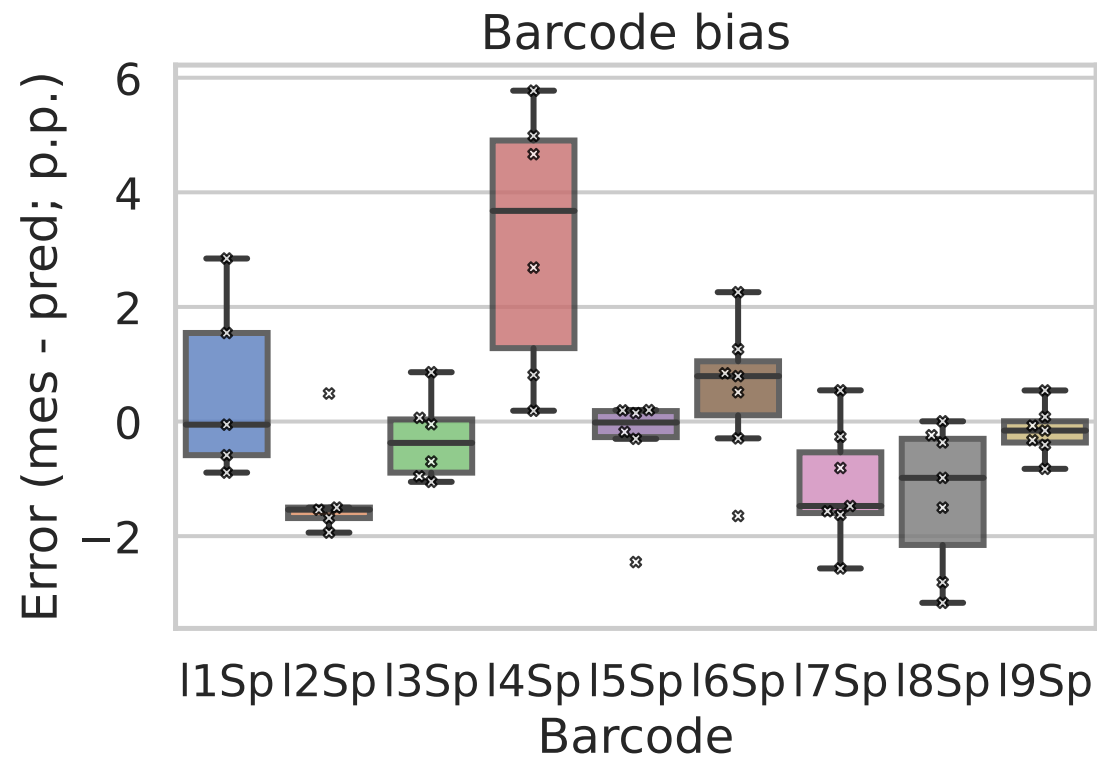
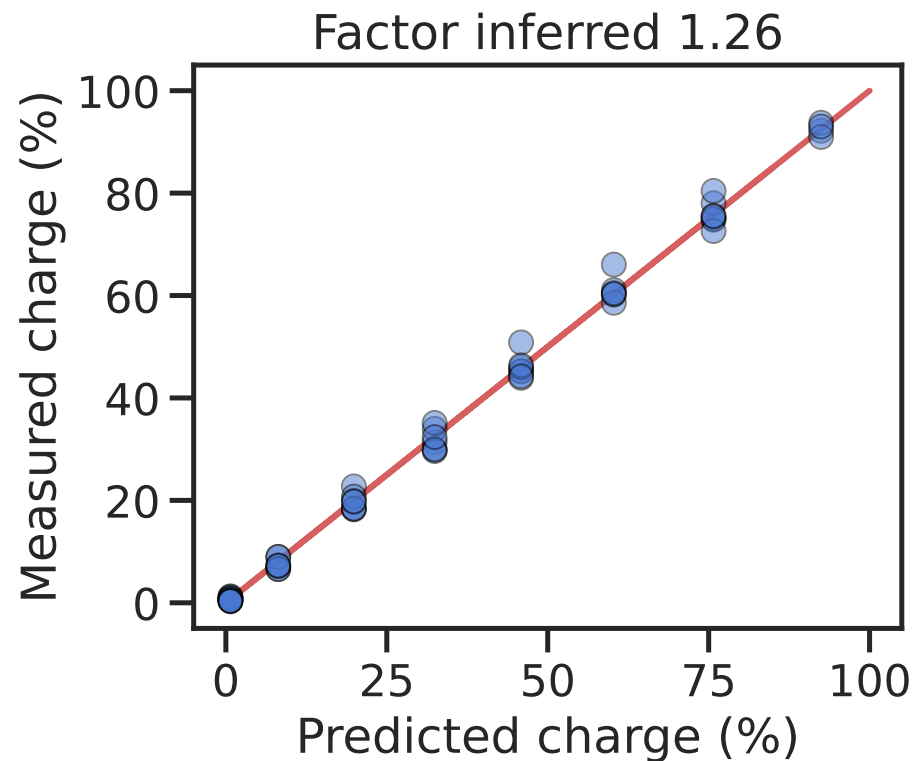
mito-Gly-TCC



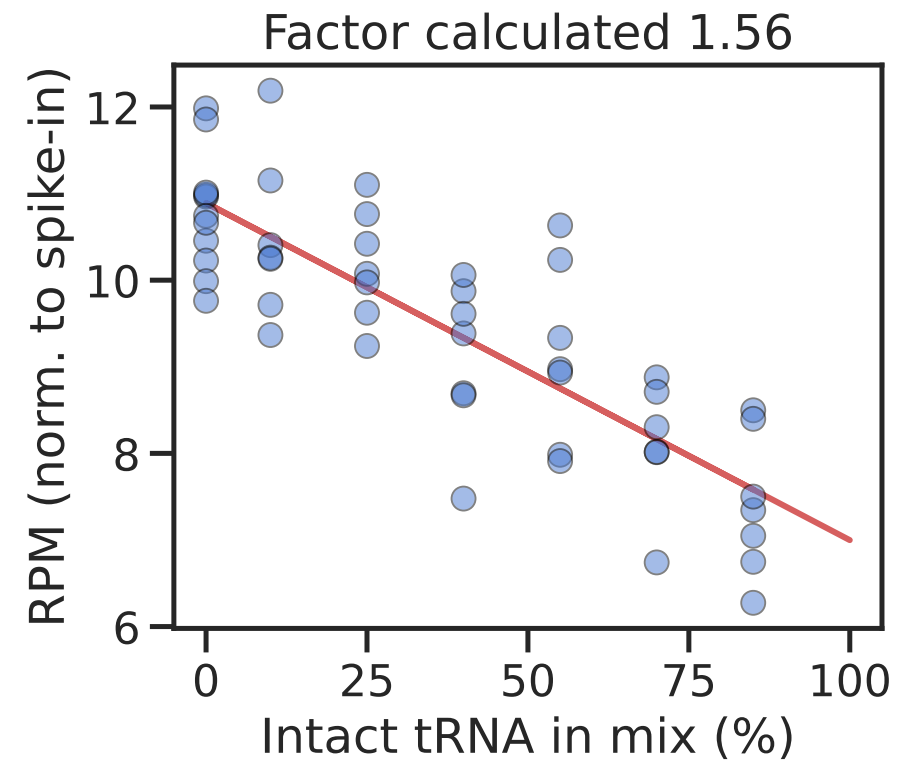
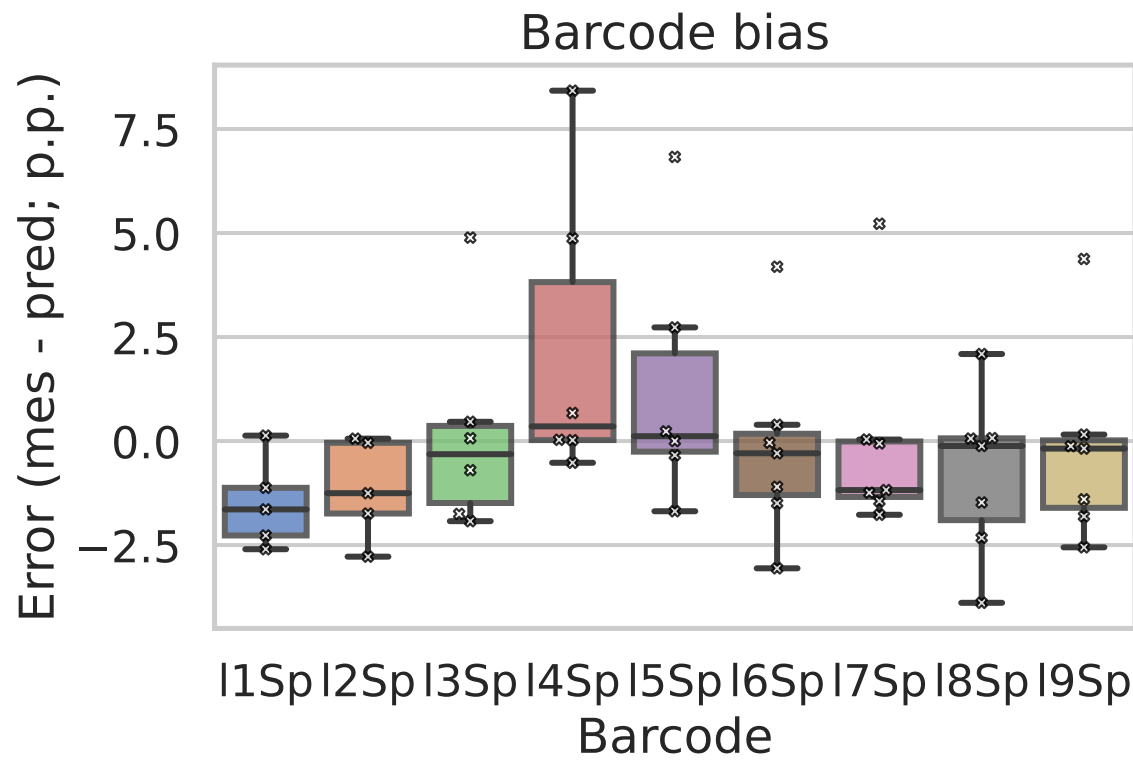
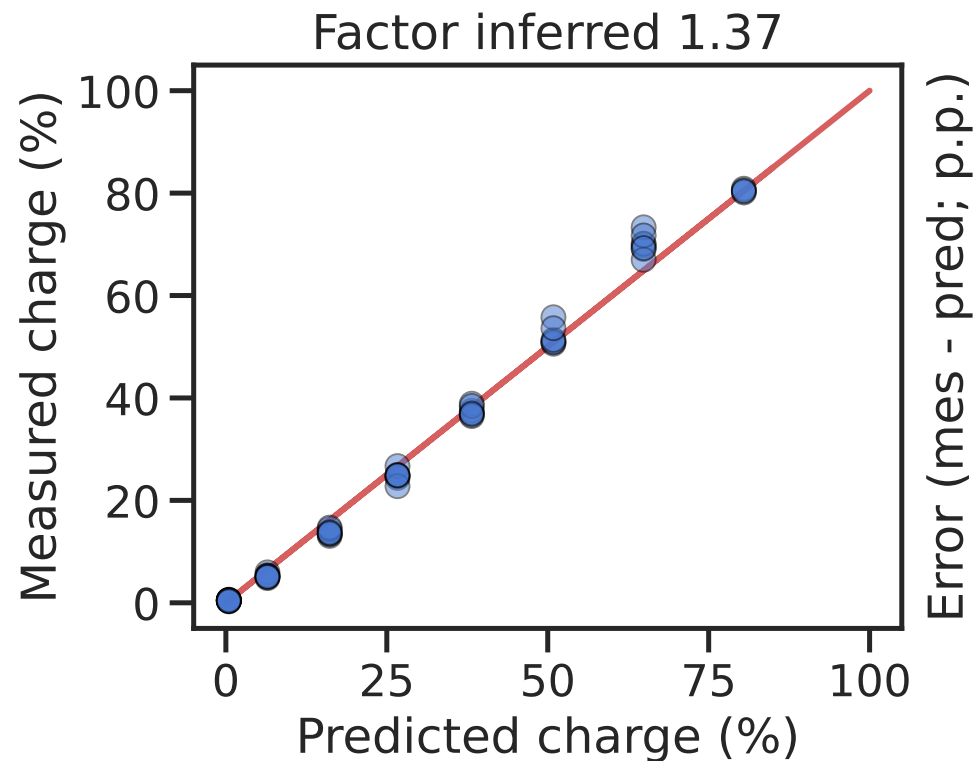
Arg-ACG-2-1



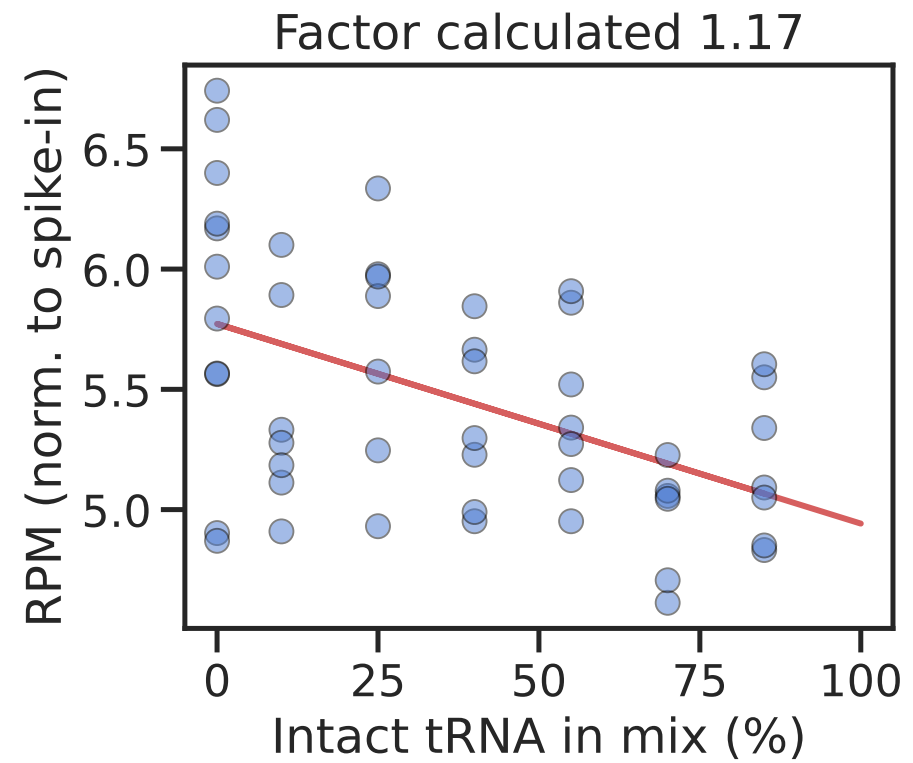
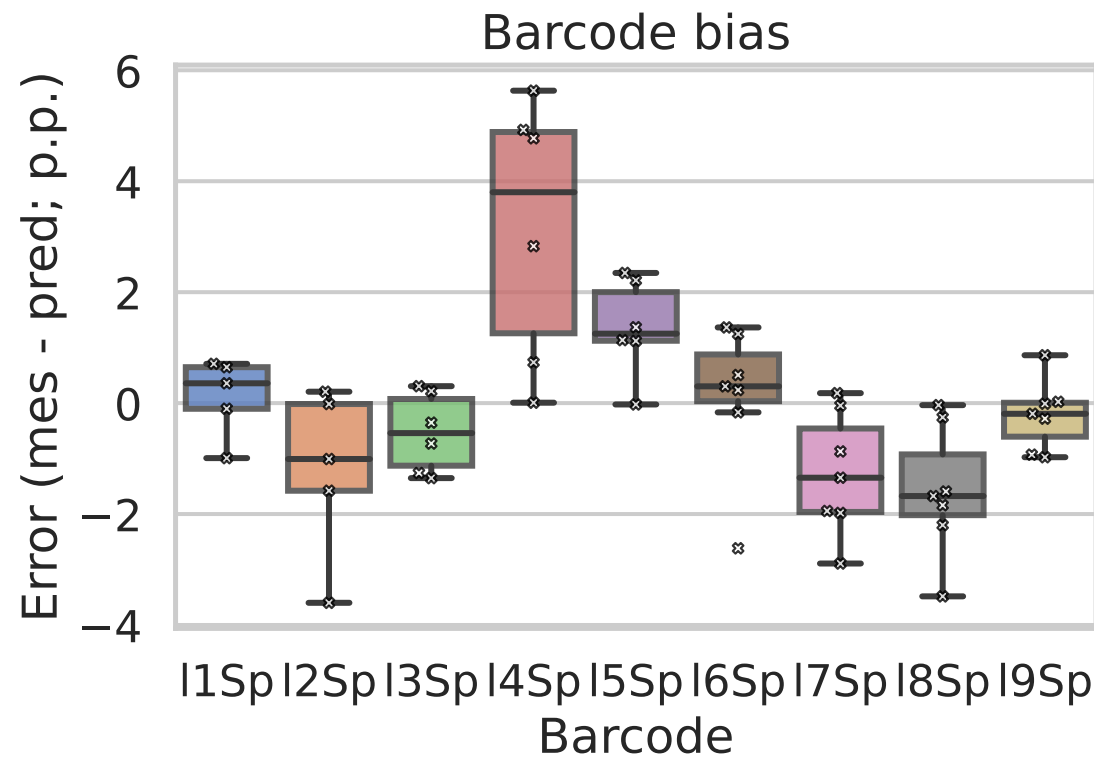
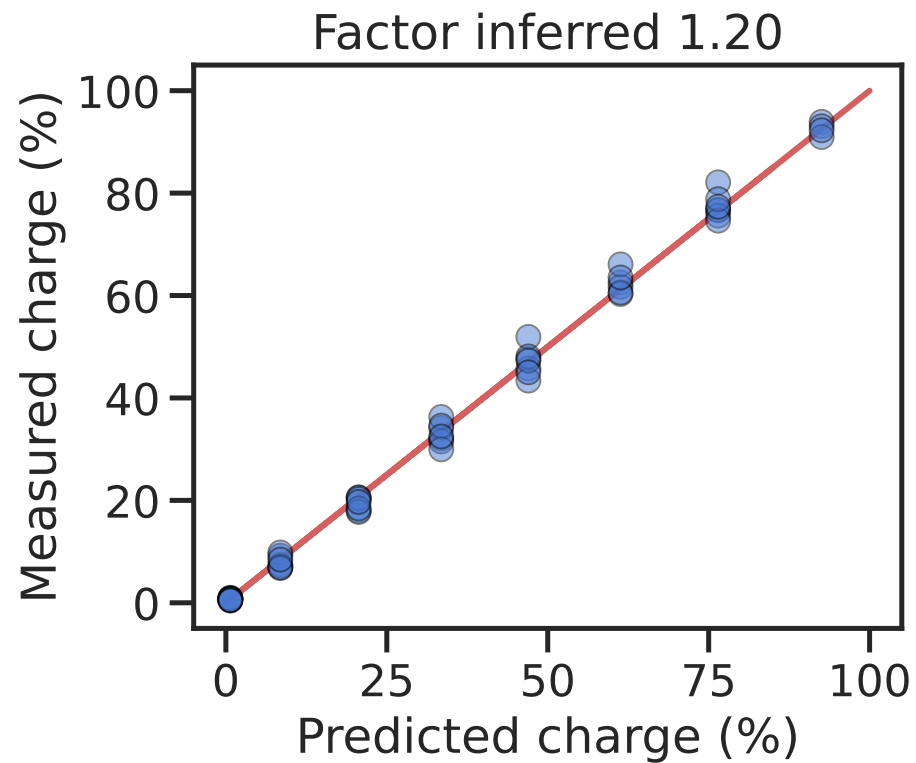
Thr-AGT-3-1



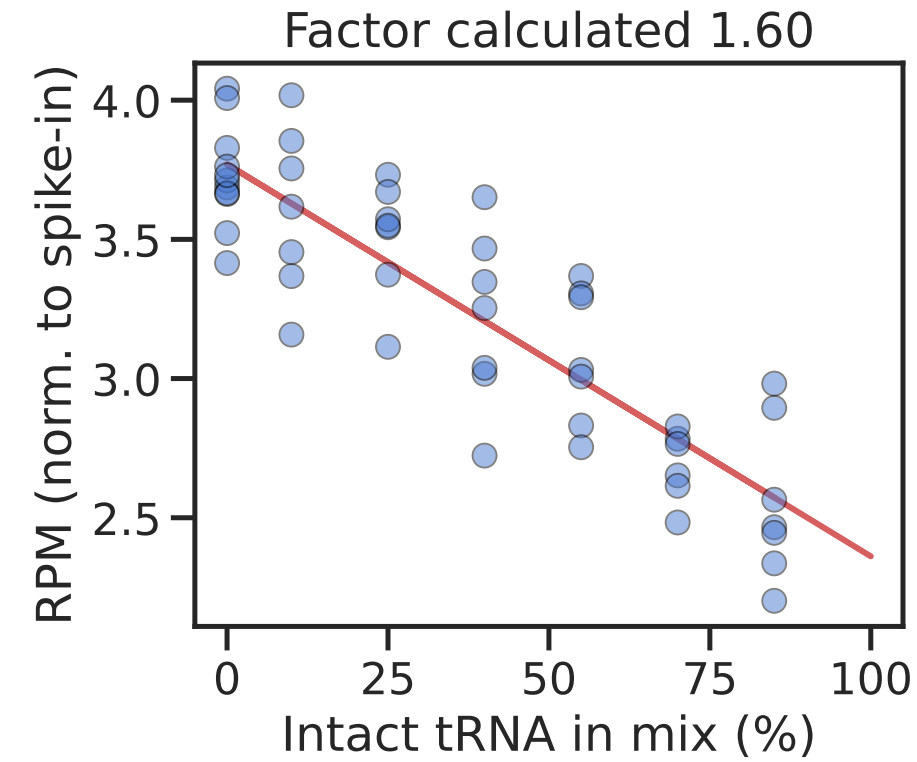
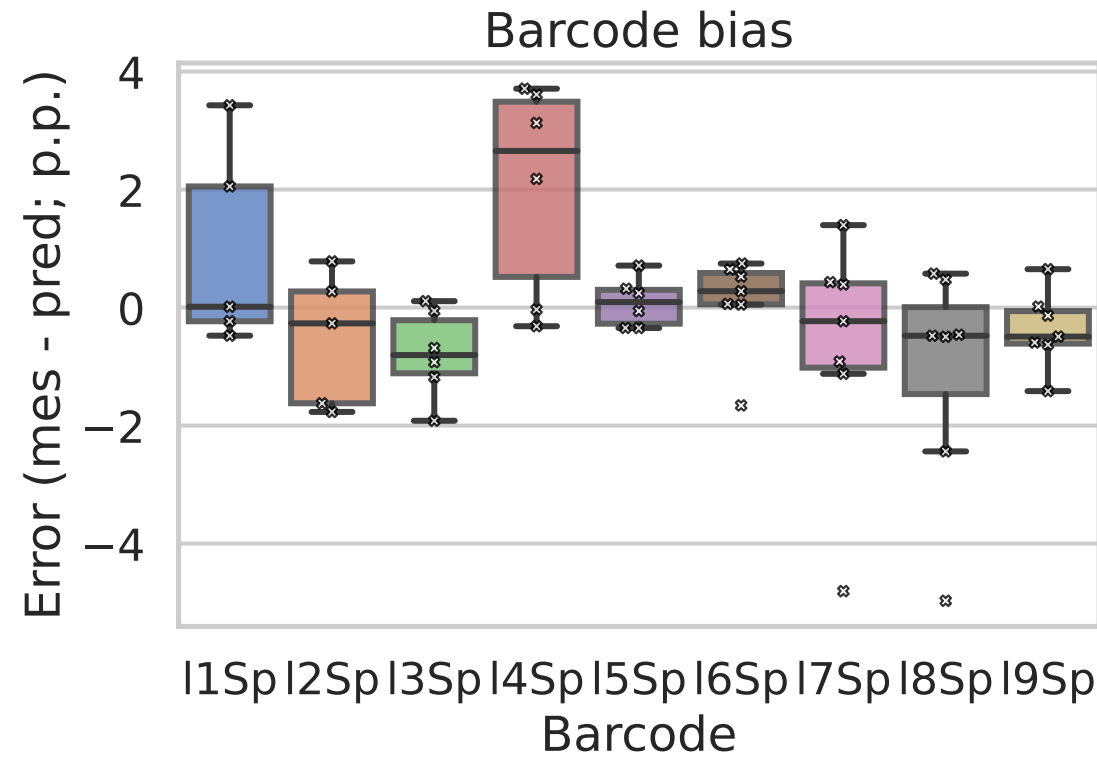
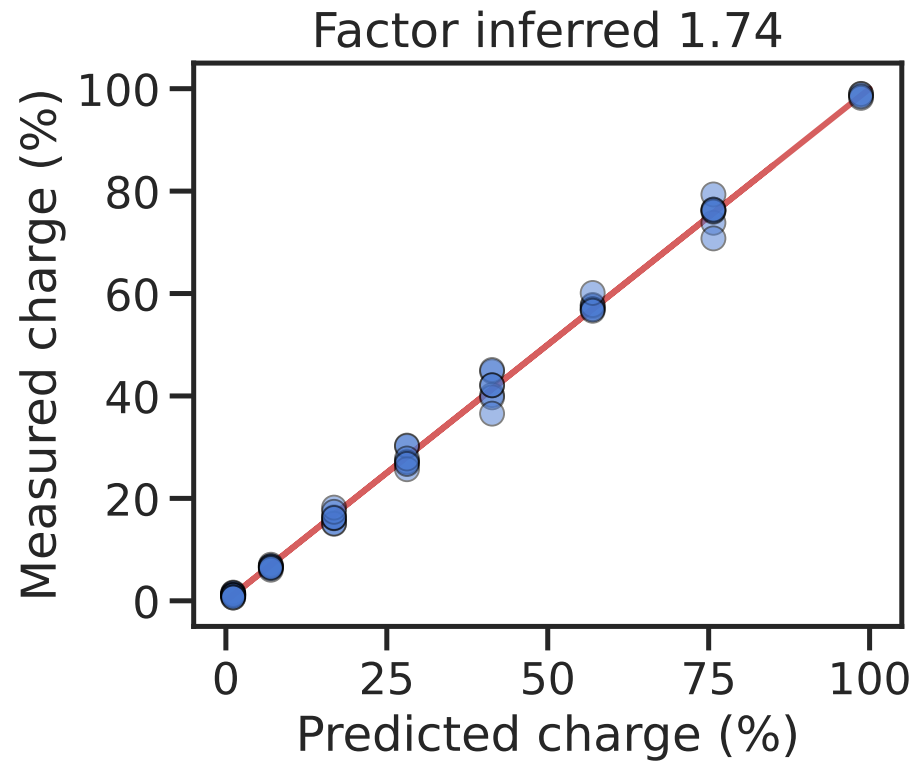
Ser-CGA-1-1



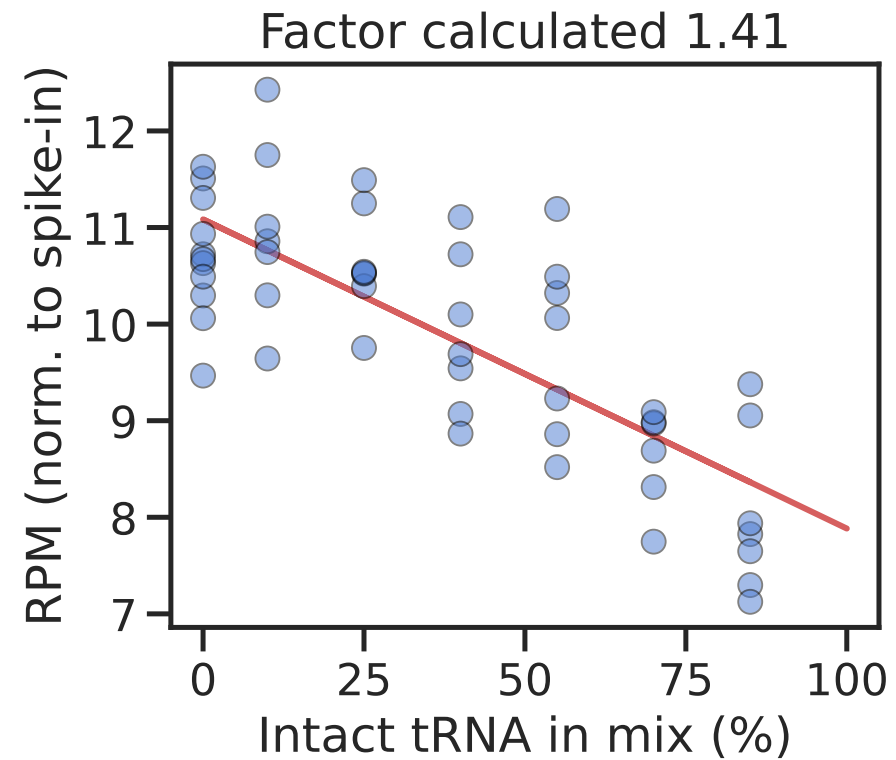
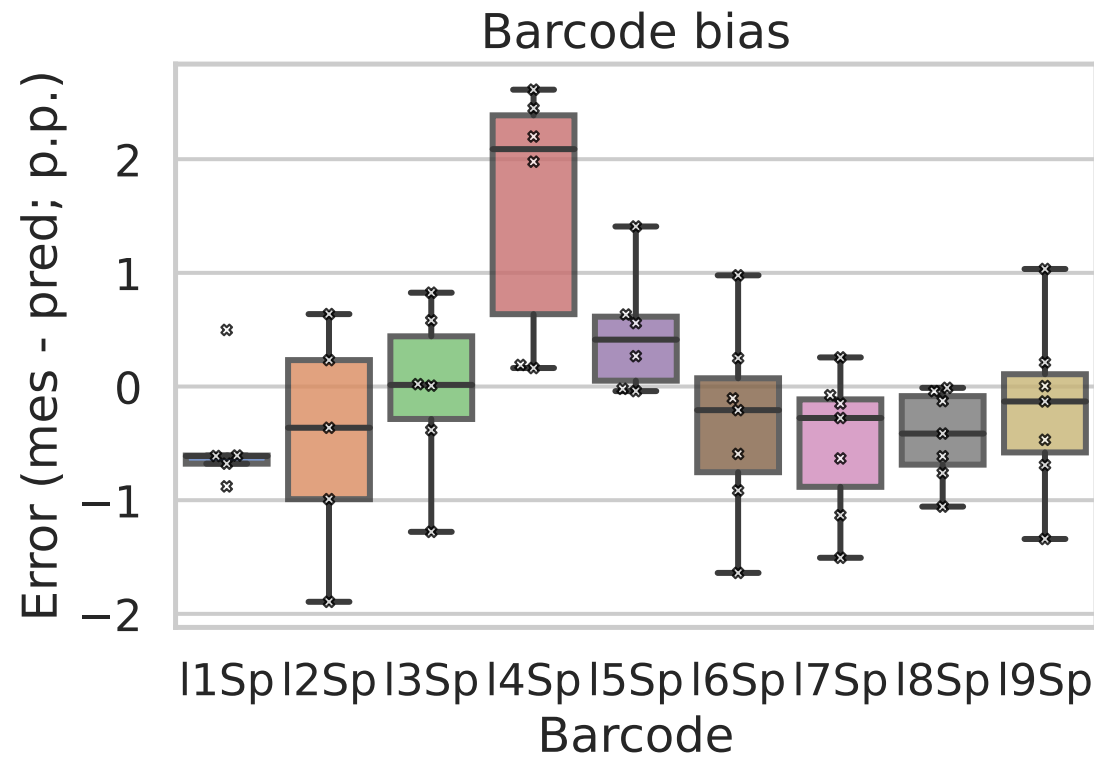
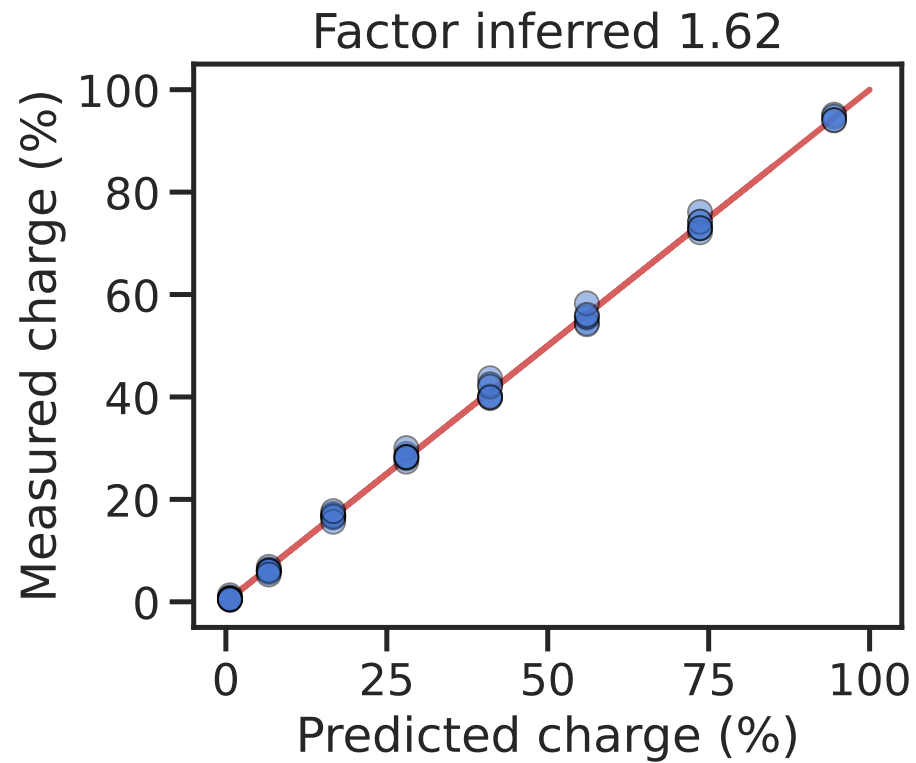
Thr-TGT-2-1



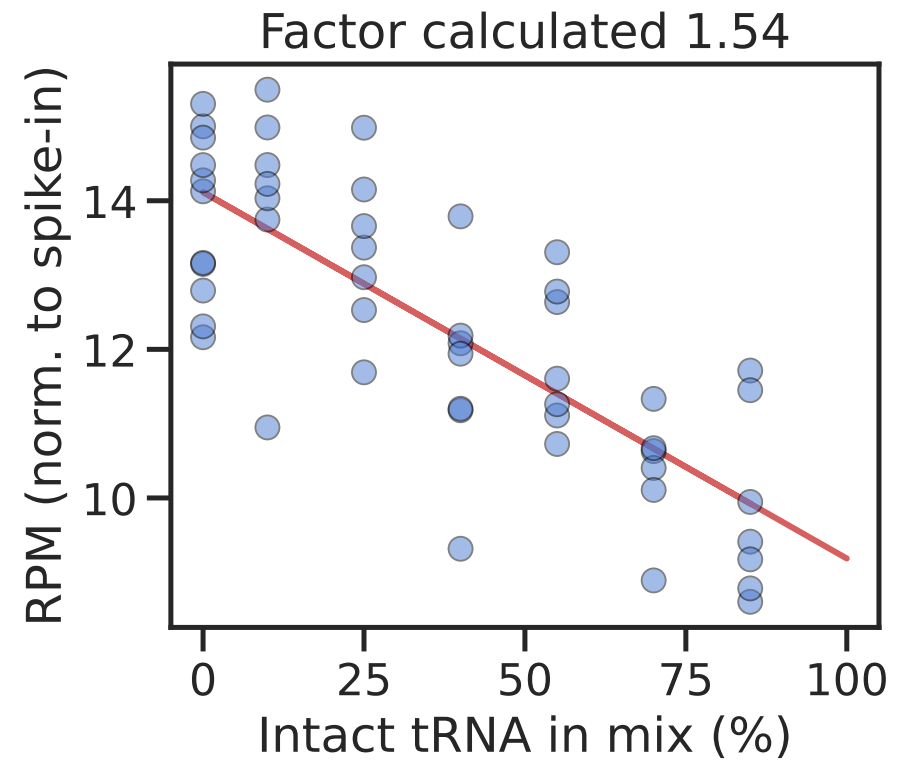
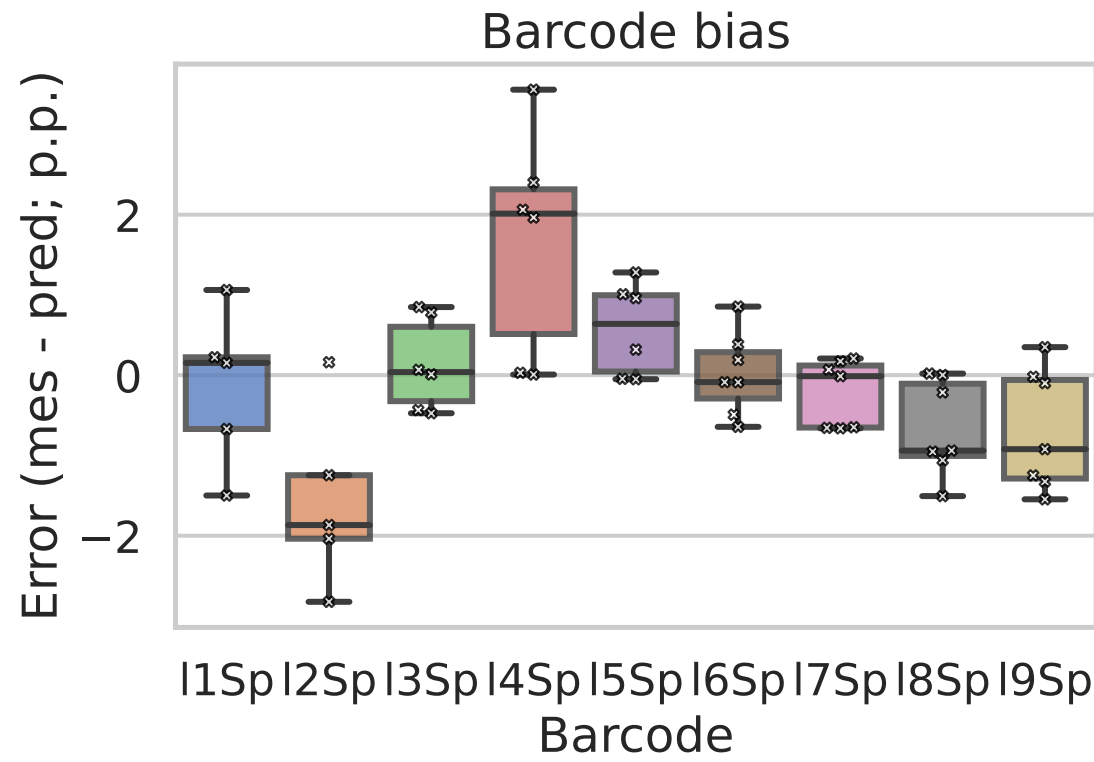
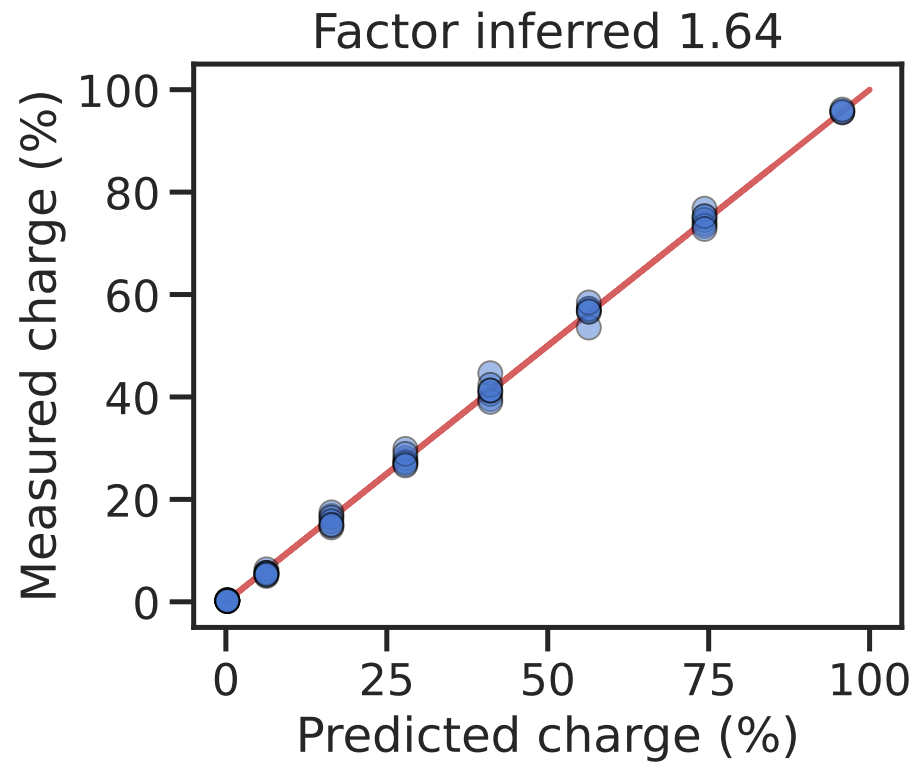
Val-AAC-3-1



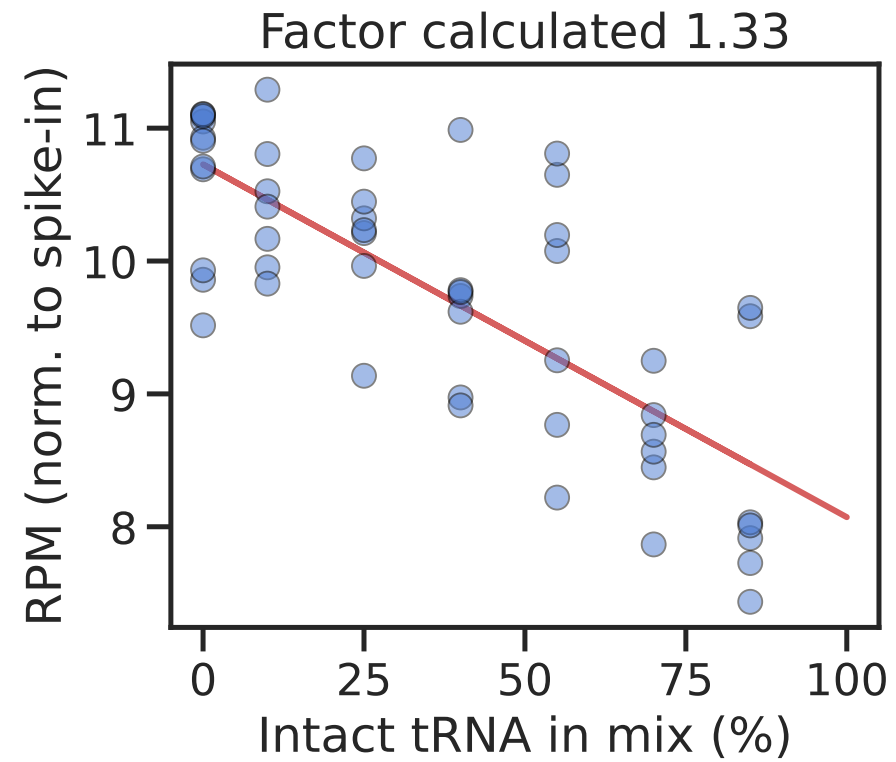
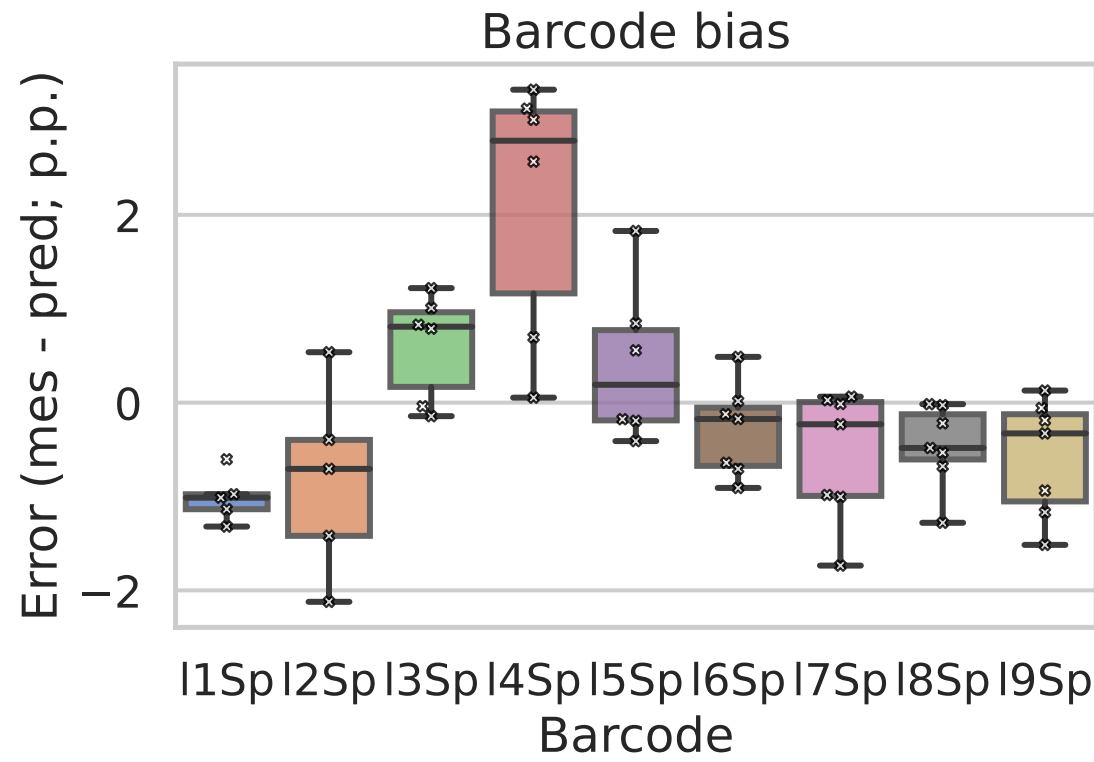
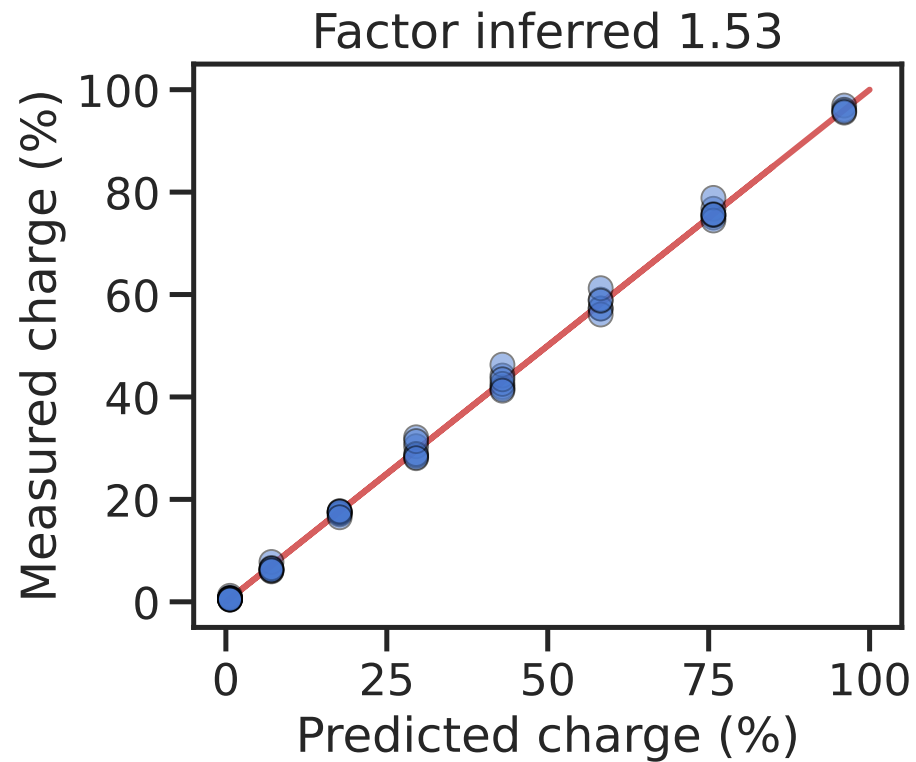
Arg-CCT-4-1



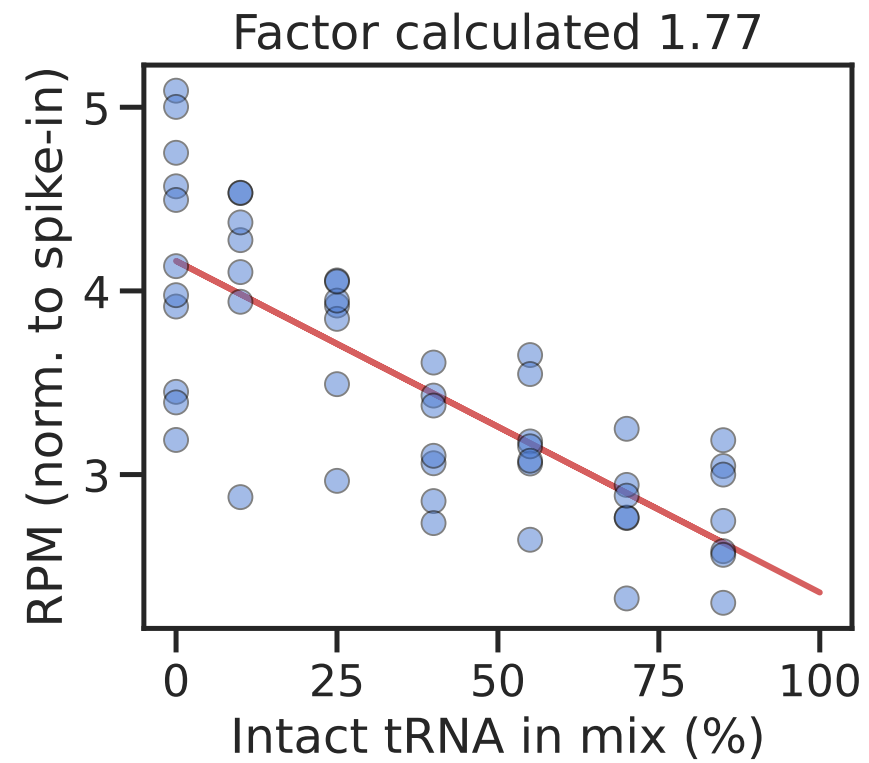
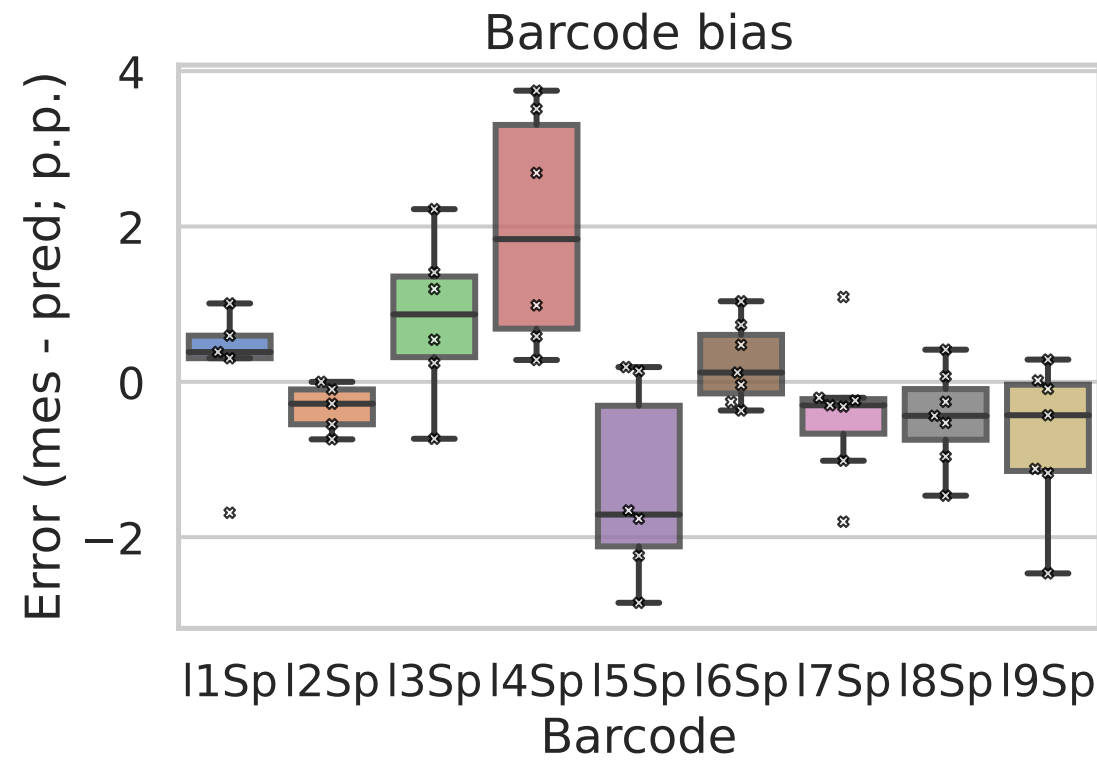
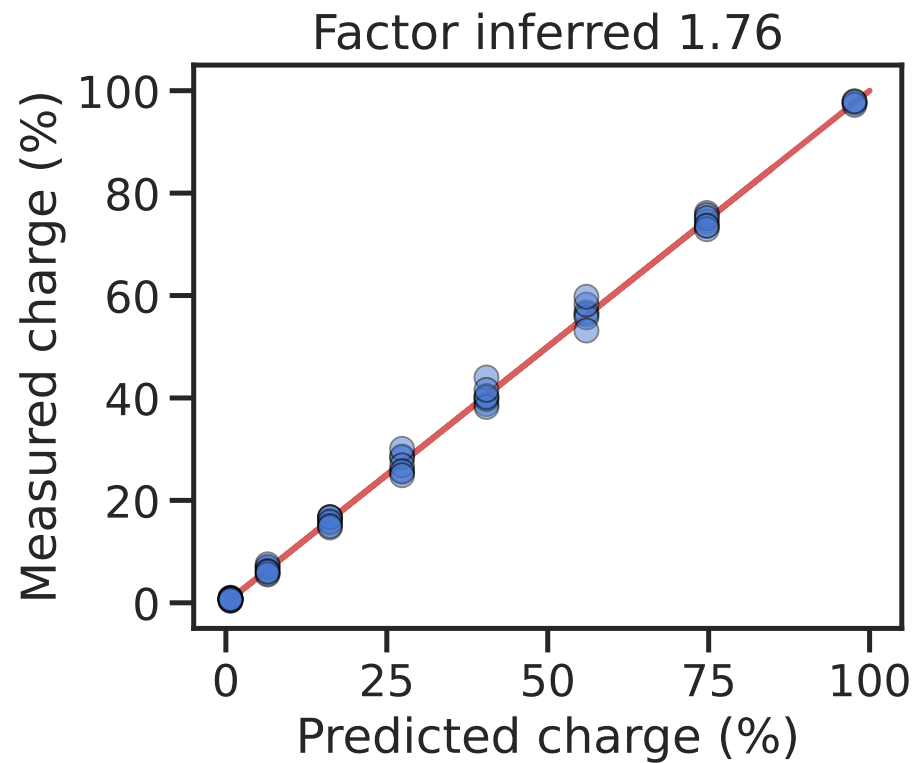
Asn-GTT-3-1



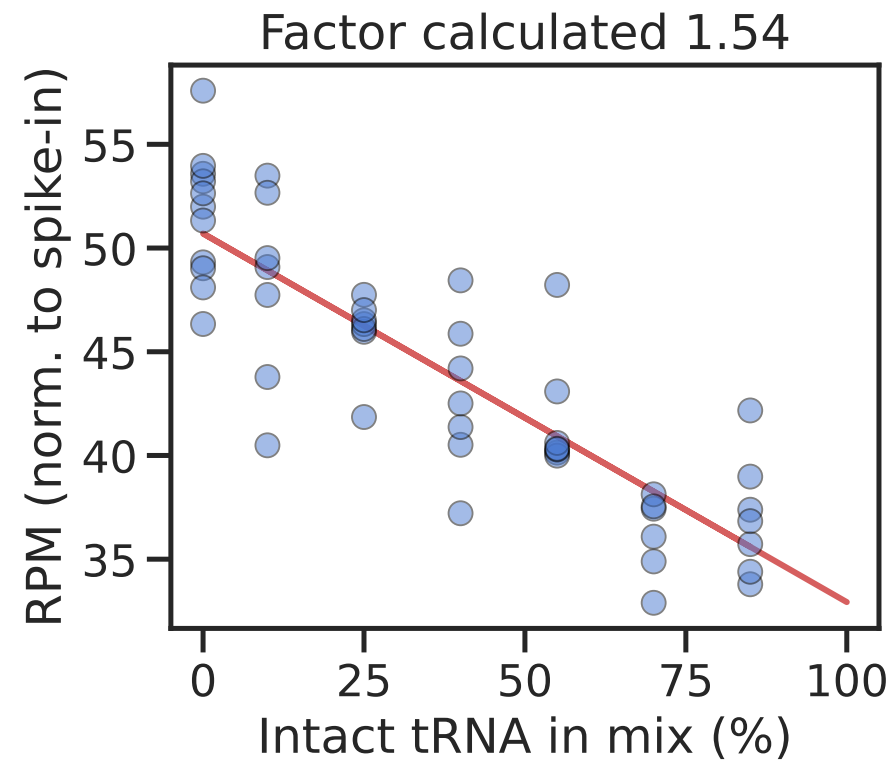
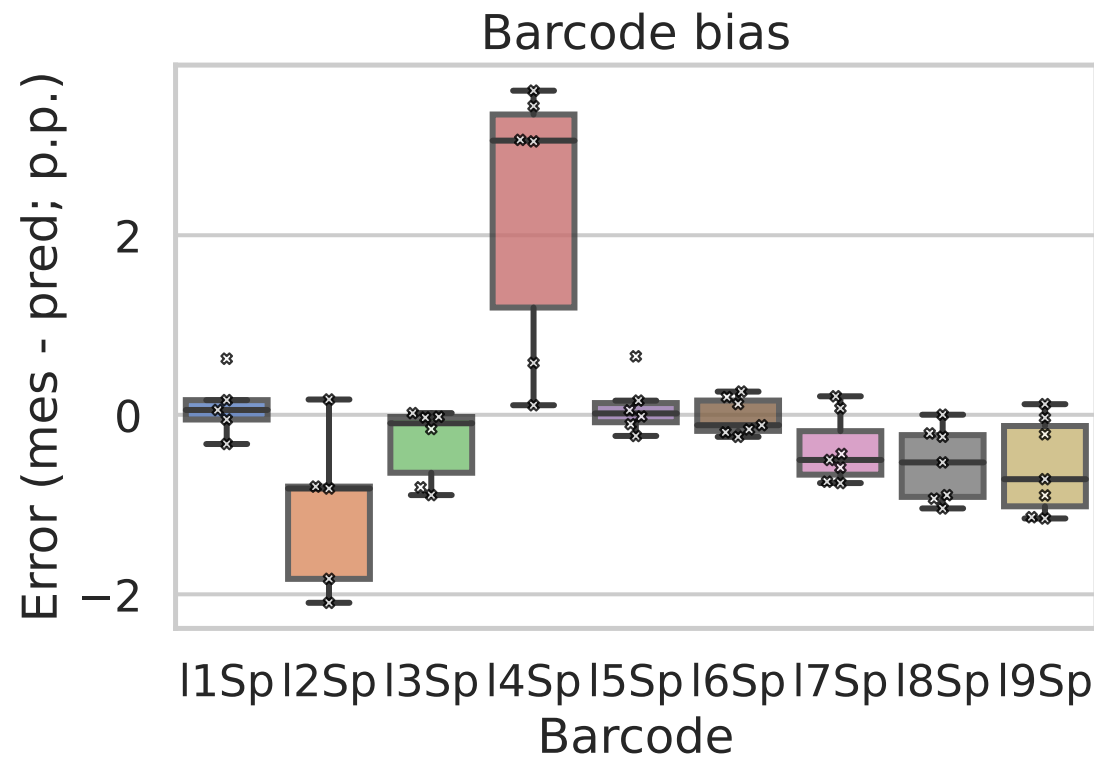
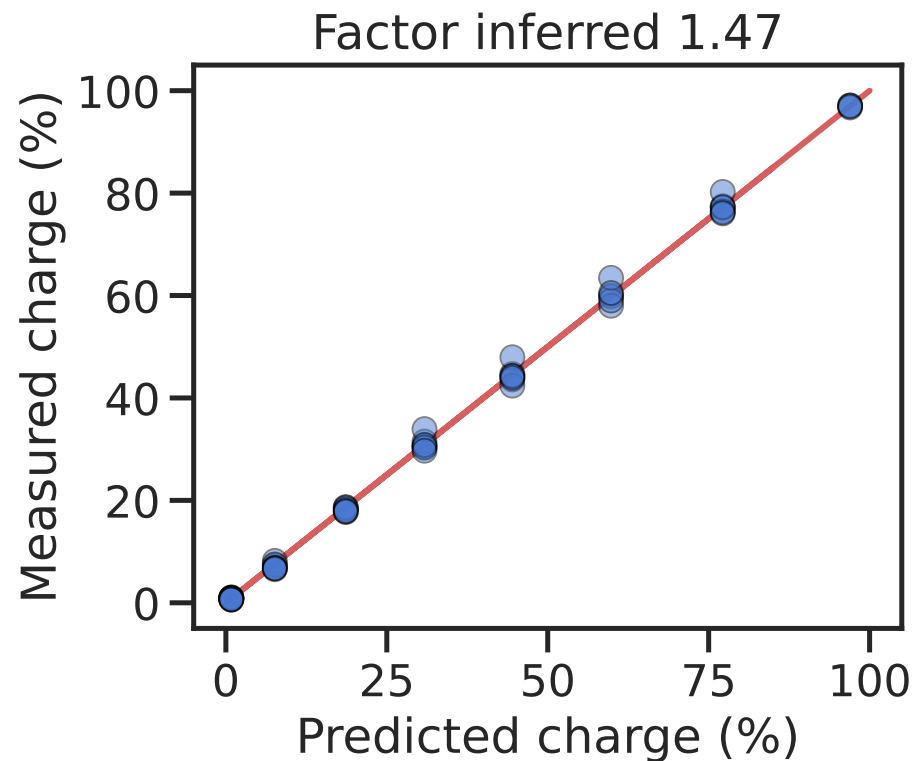
Met-CAT-6-1



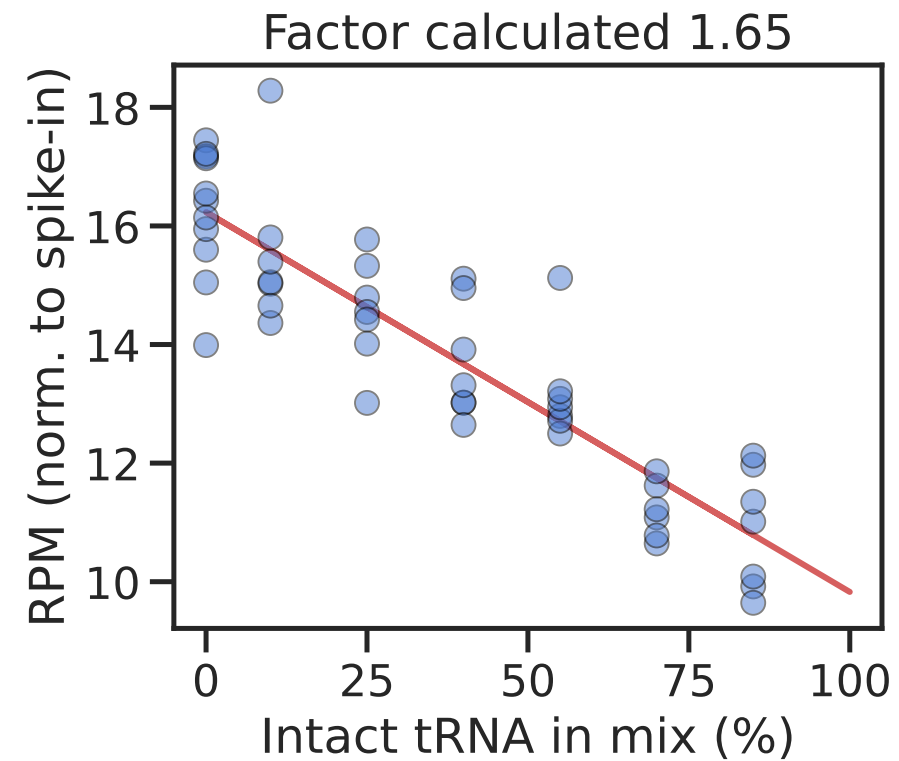
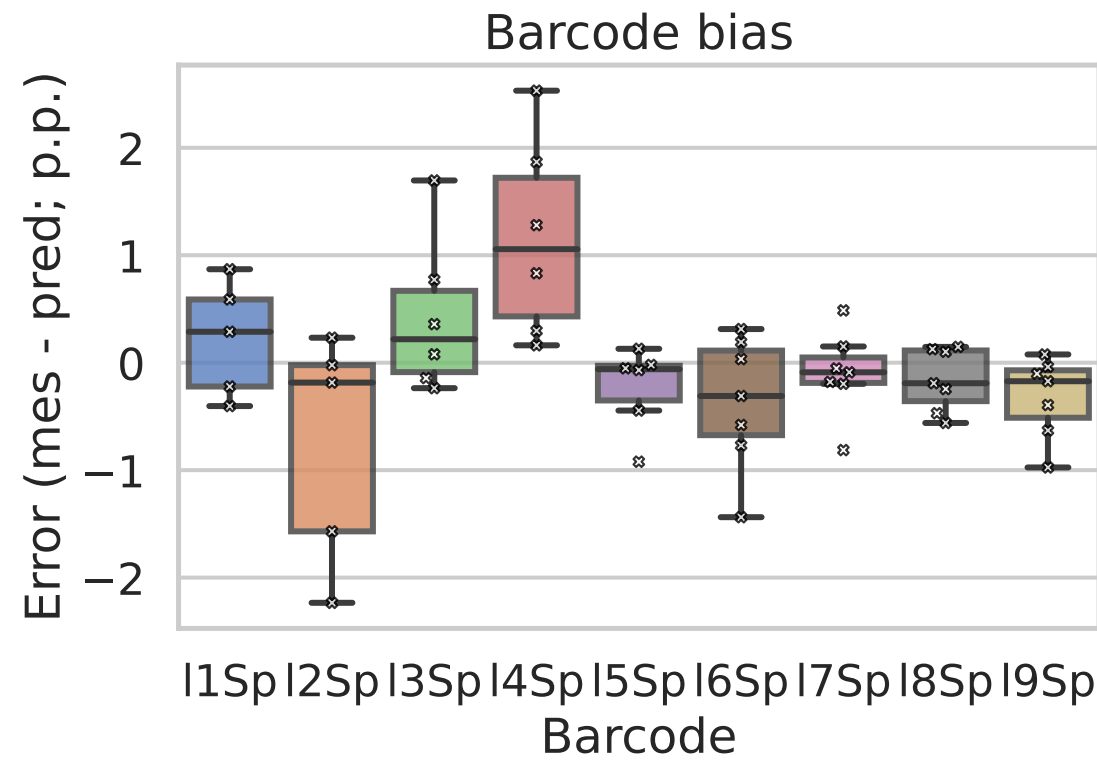
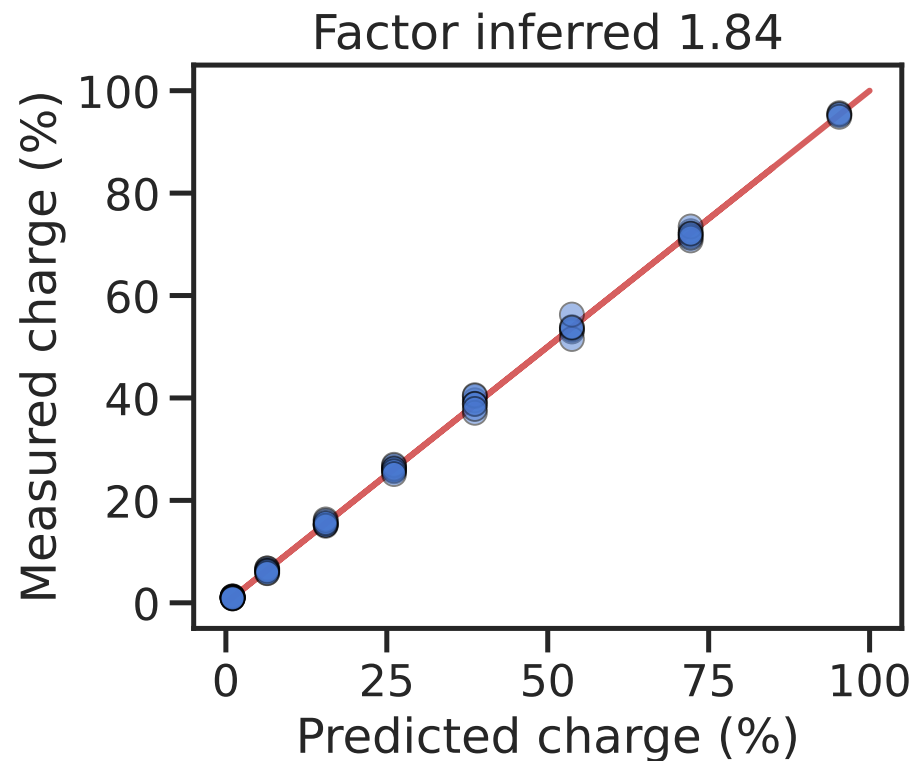
Trp-CCA-3-3



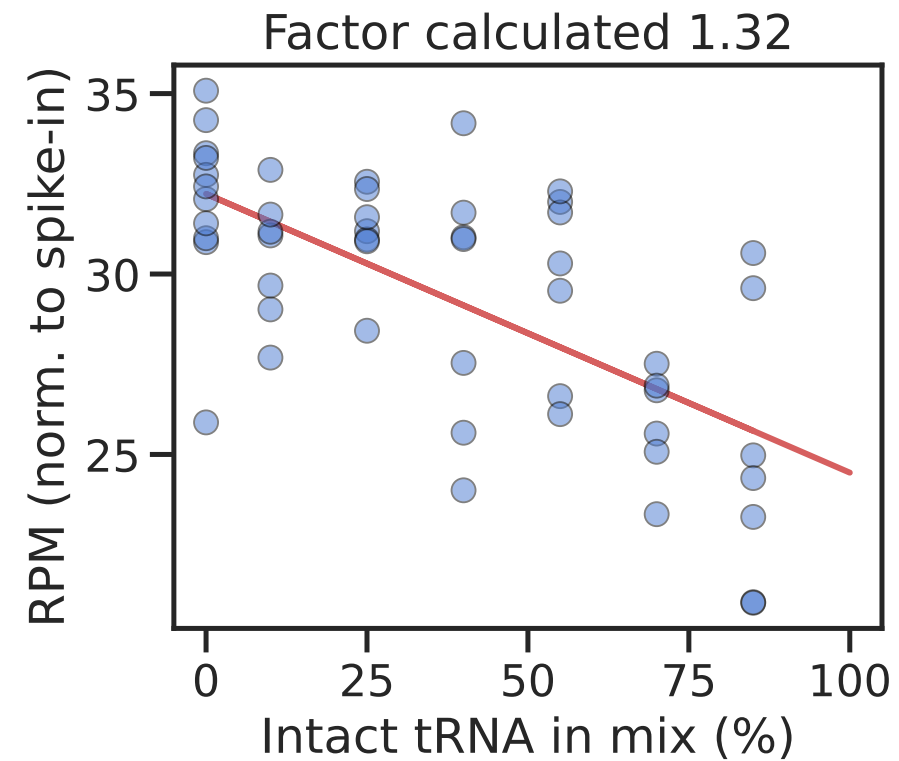
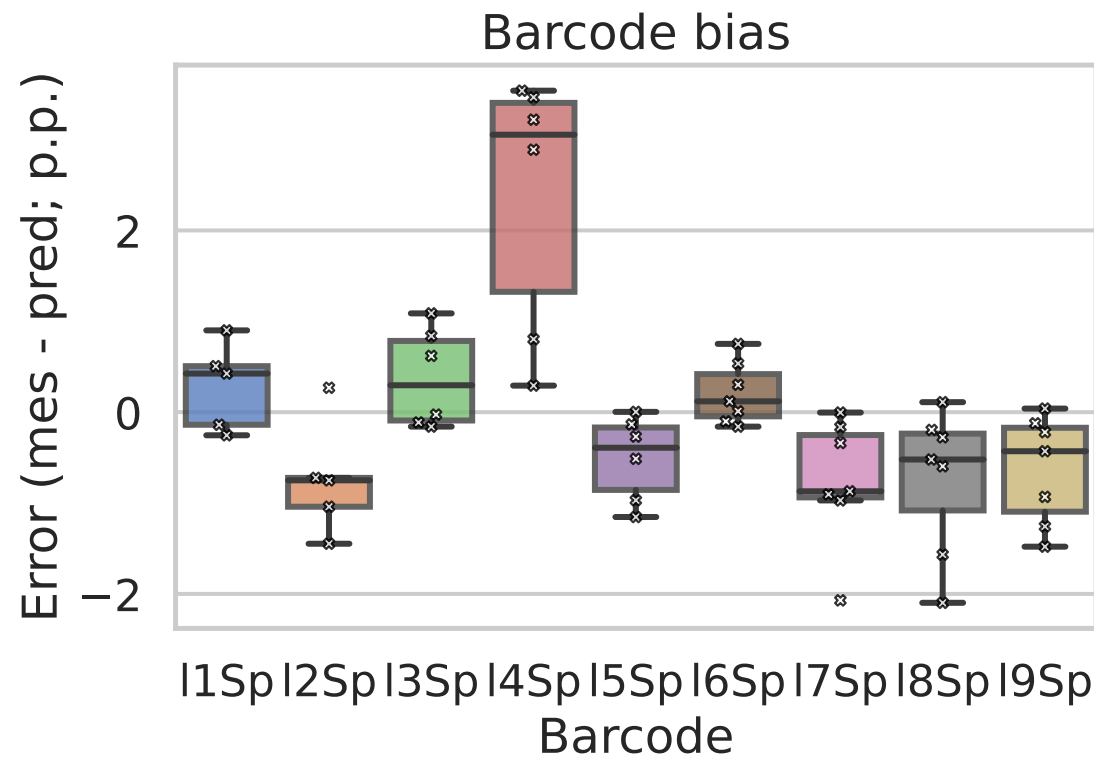
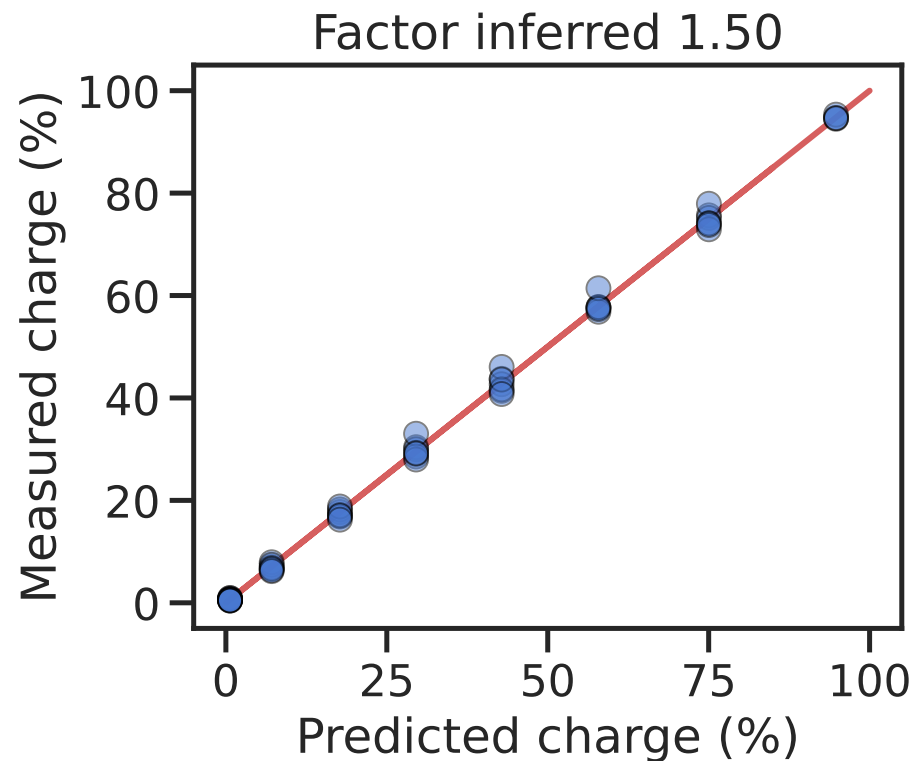
Gly-TCC-2-1



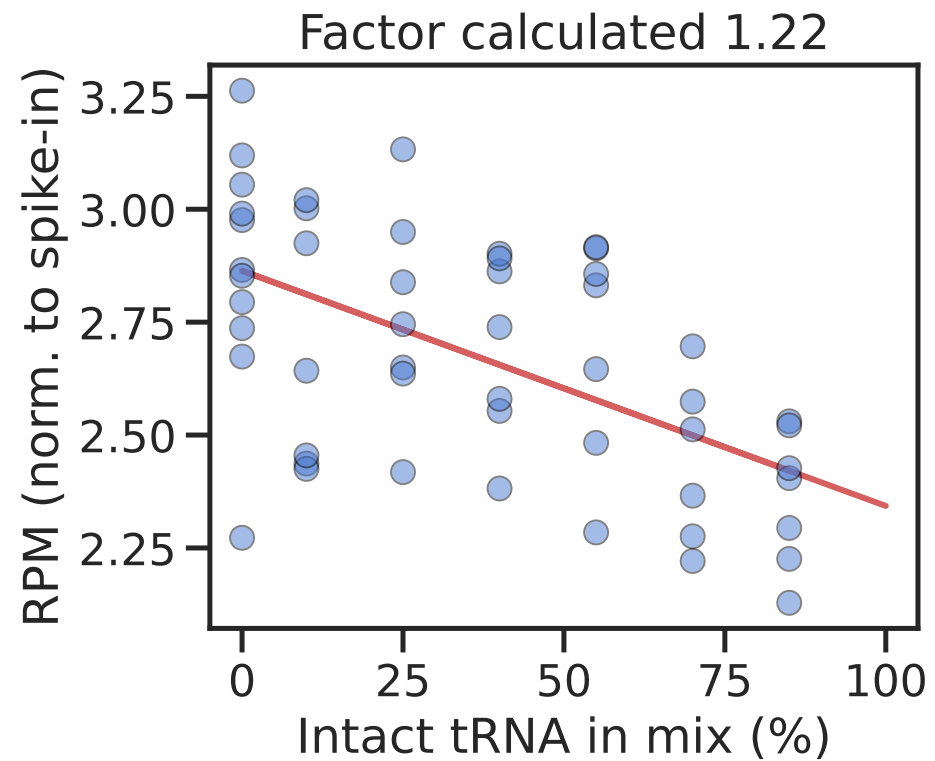
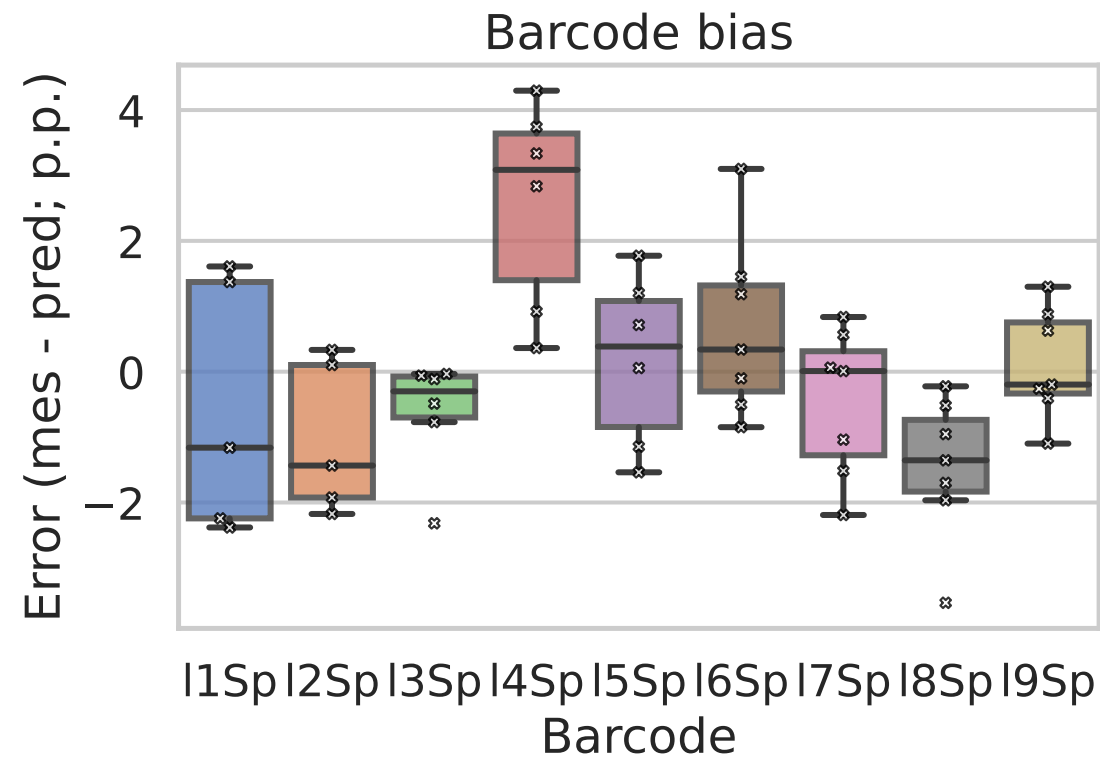
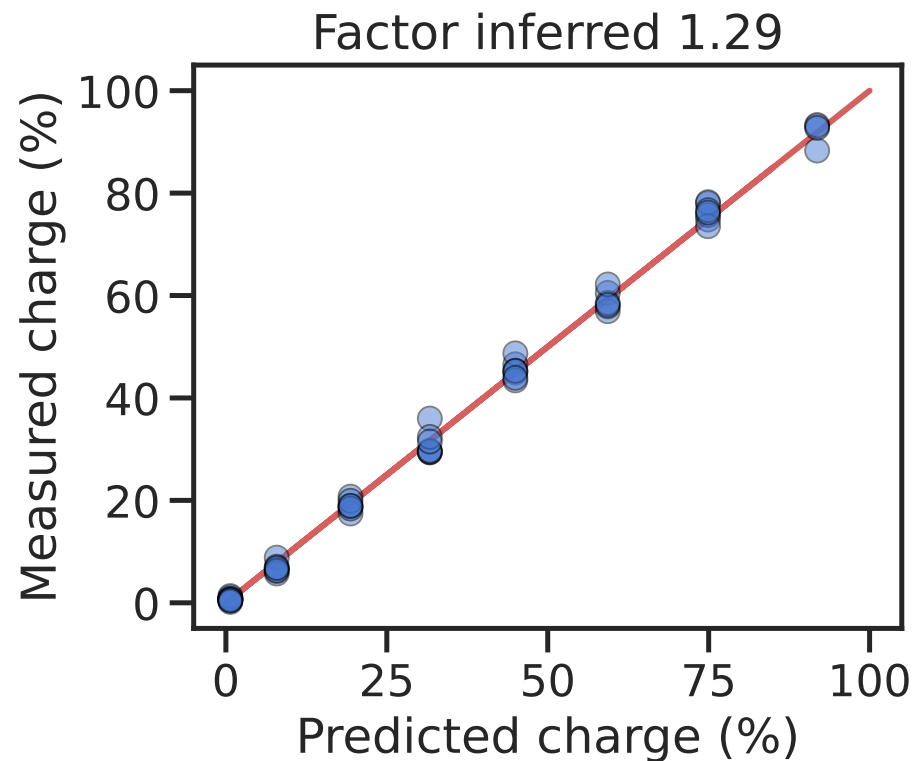
Ile-AAT-6-1



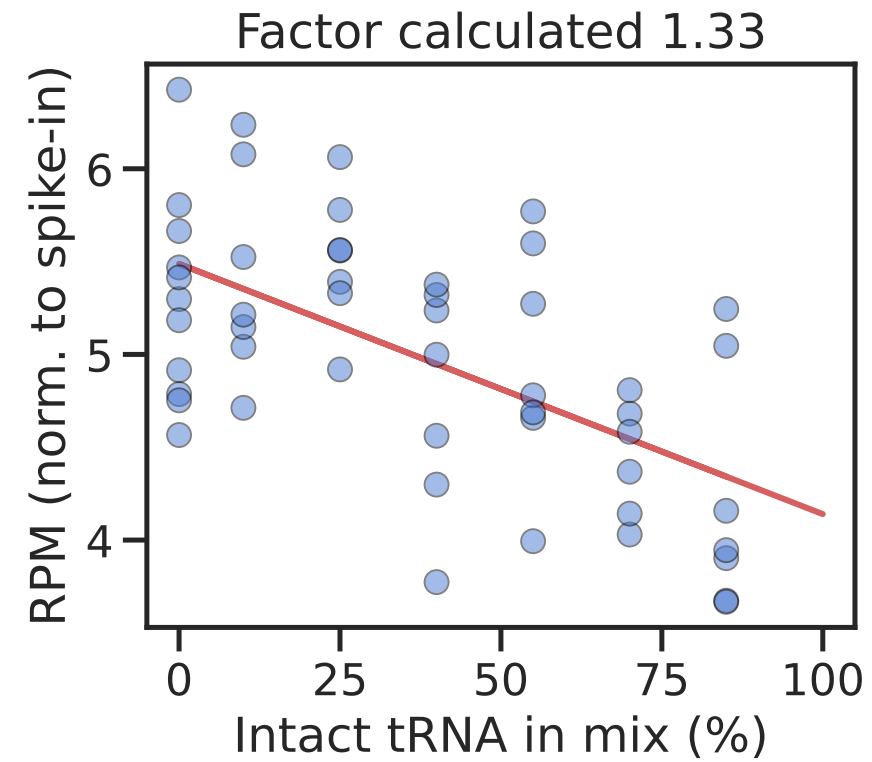
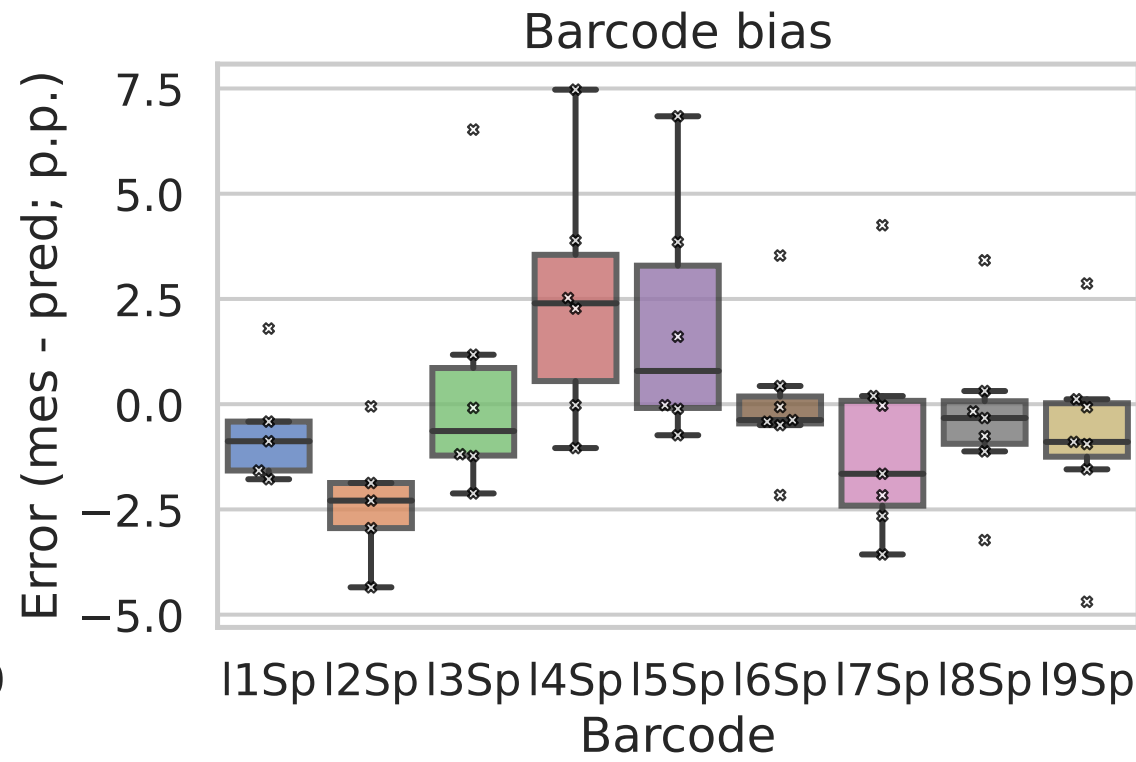
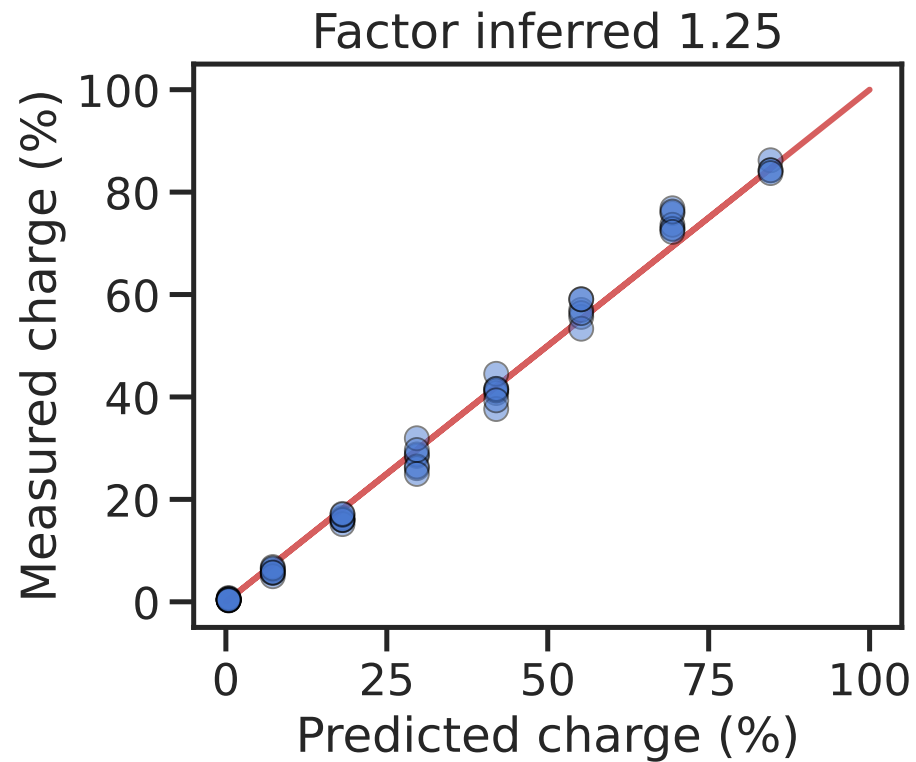
Ala-AGC-2-3



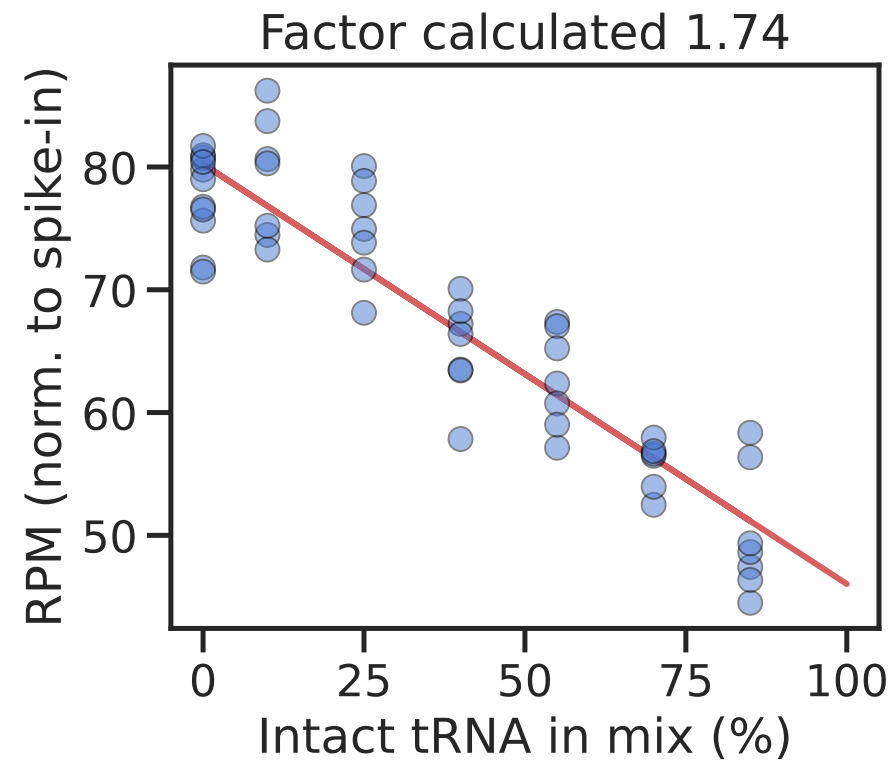
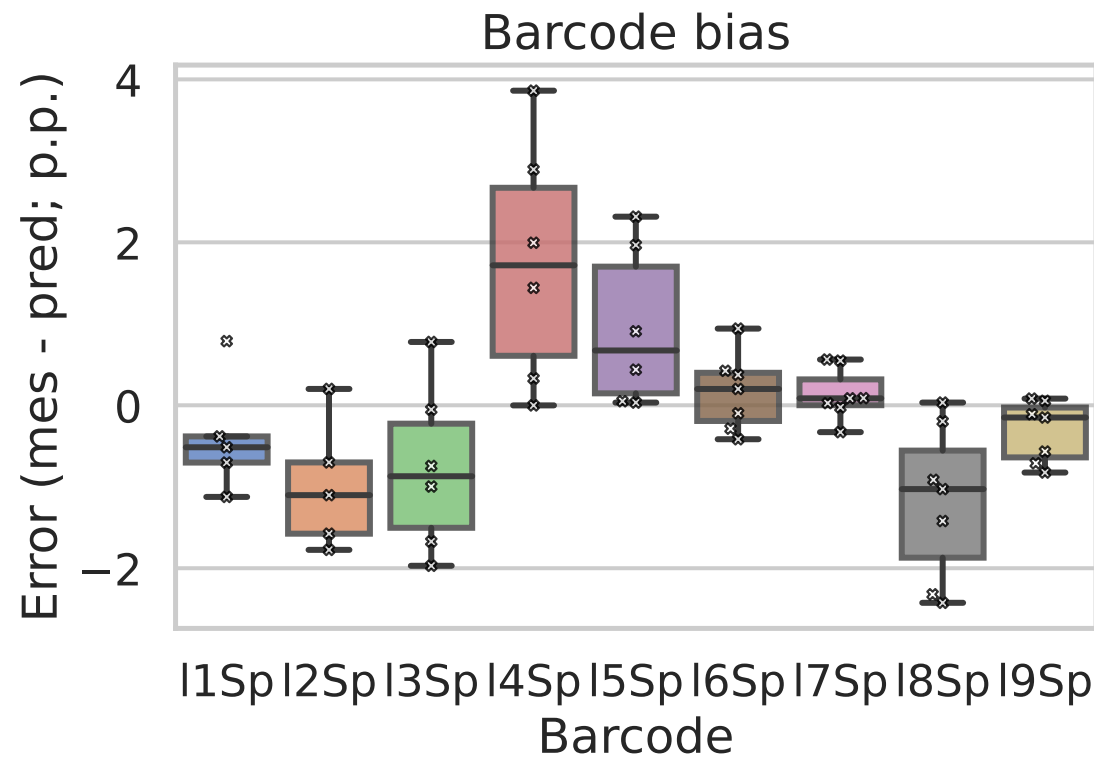
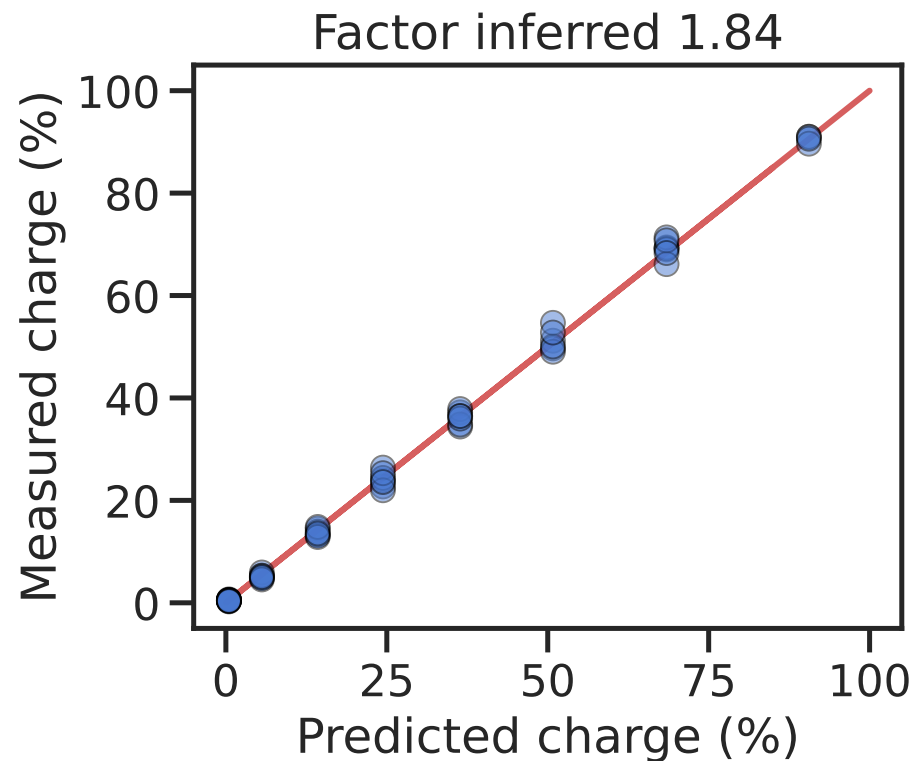
Thr-CGT-1-1



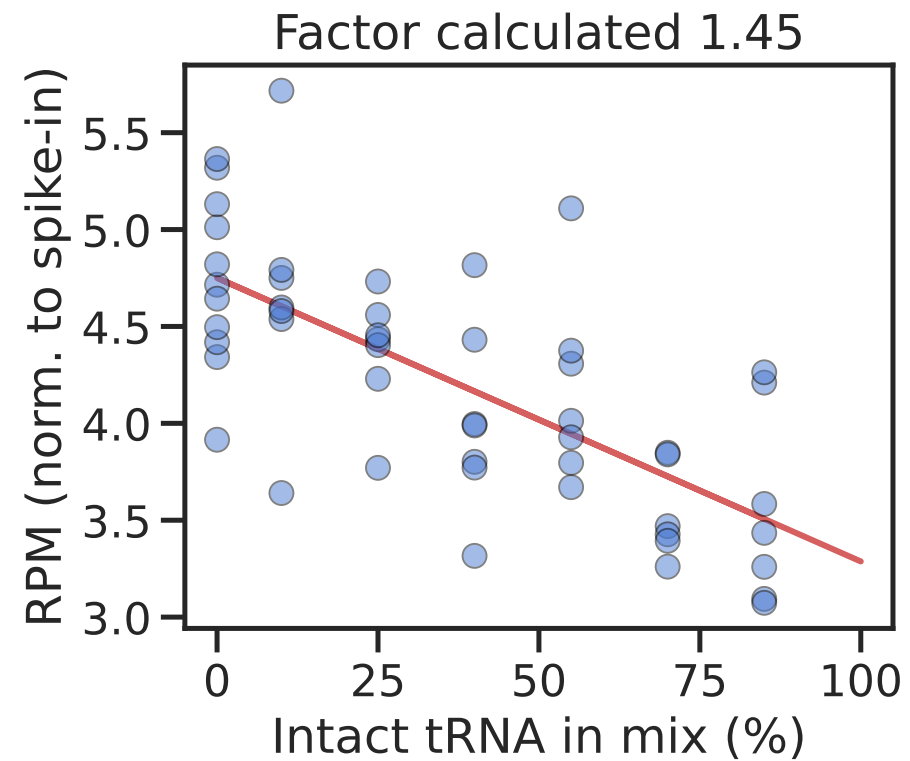
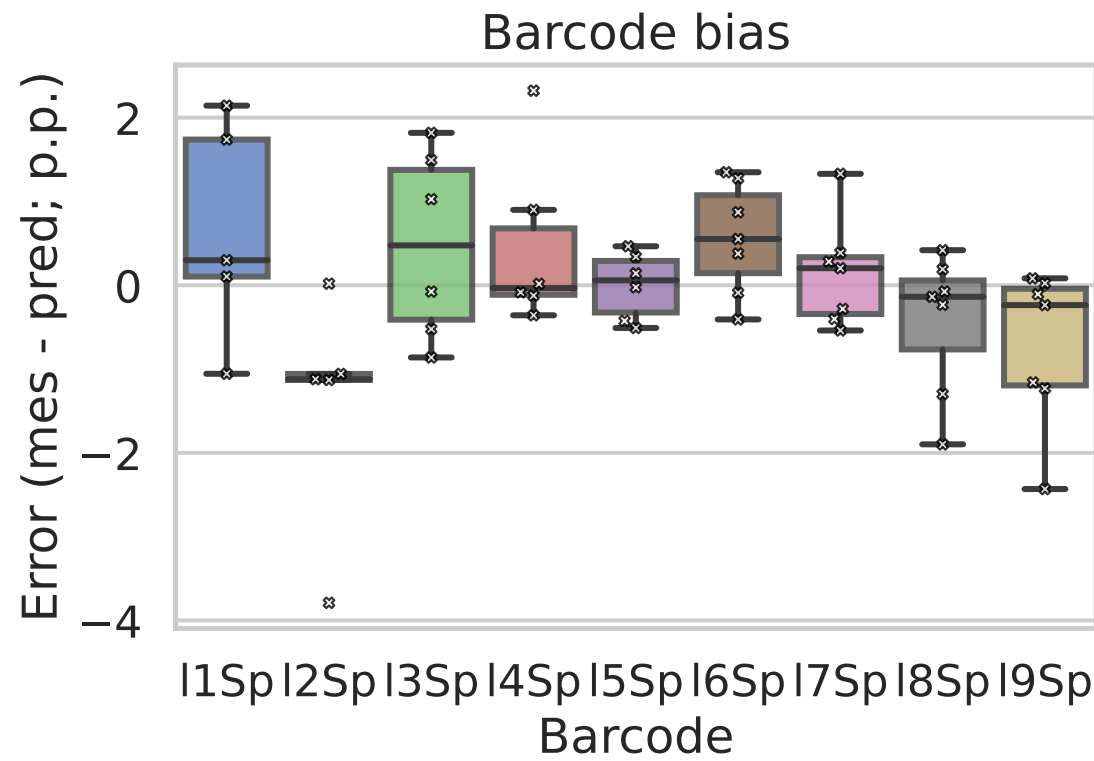
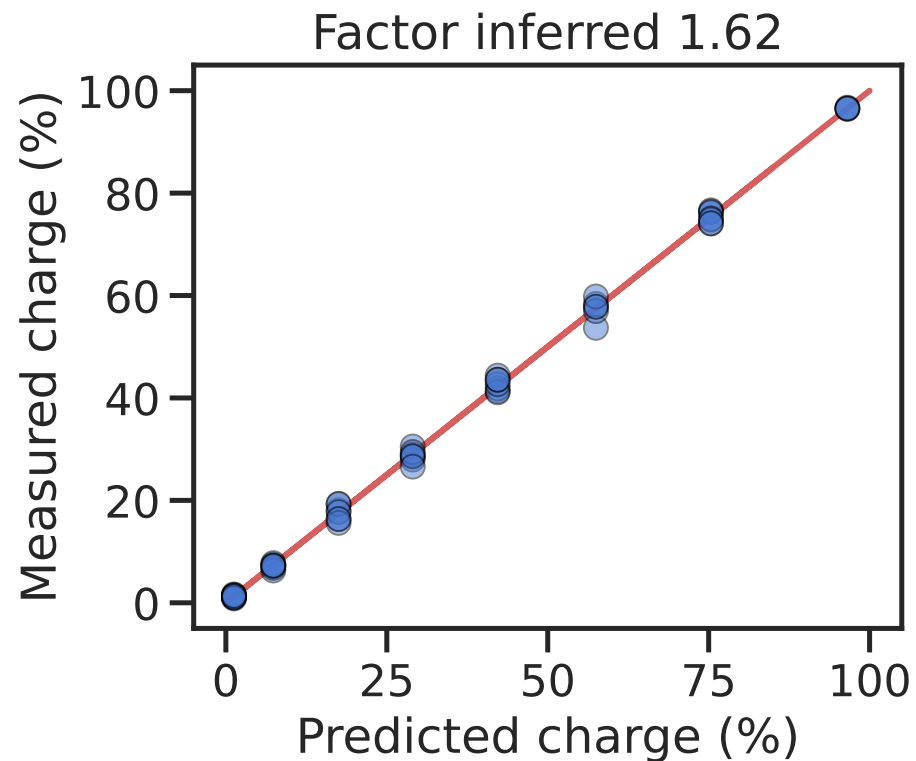
Ser-TGA-2-1



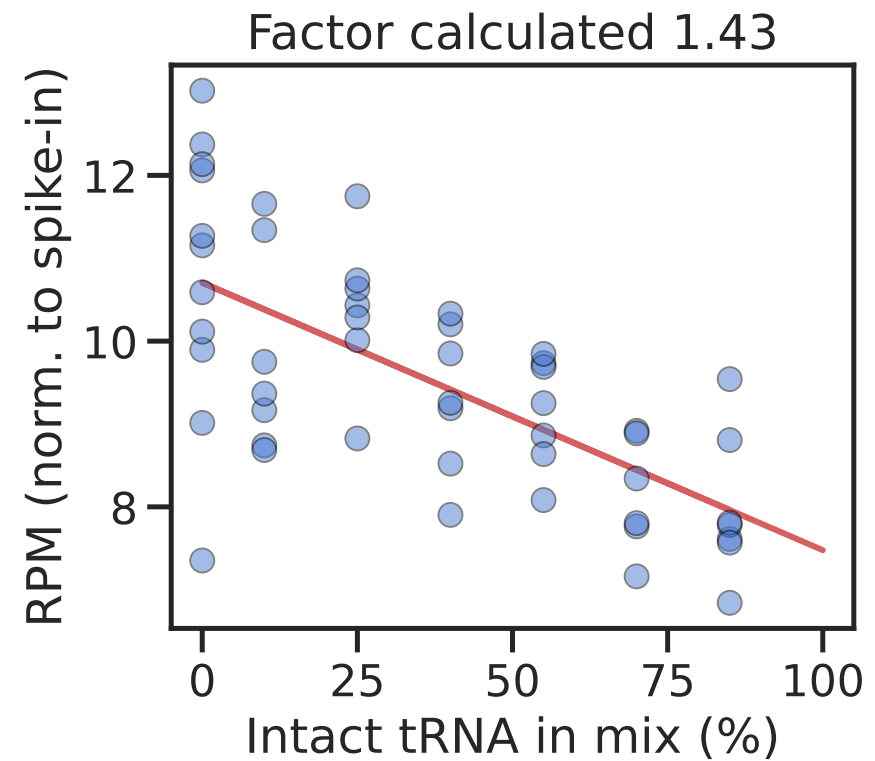
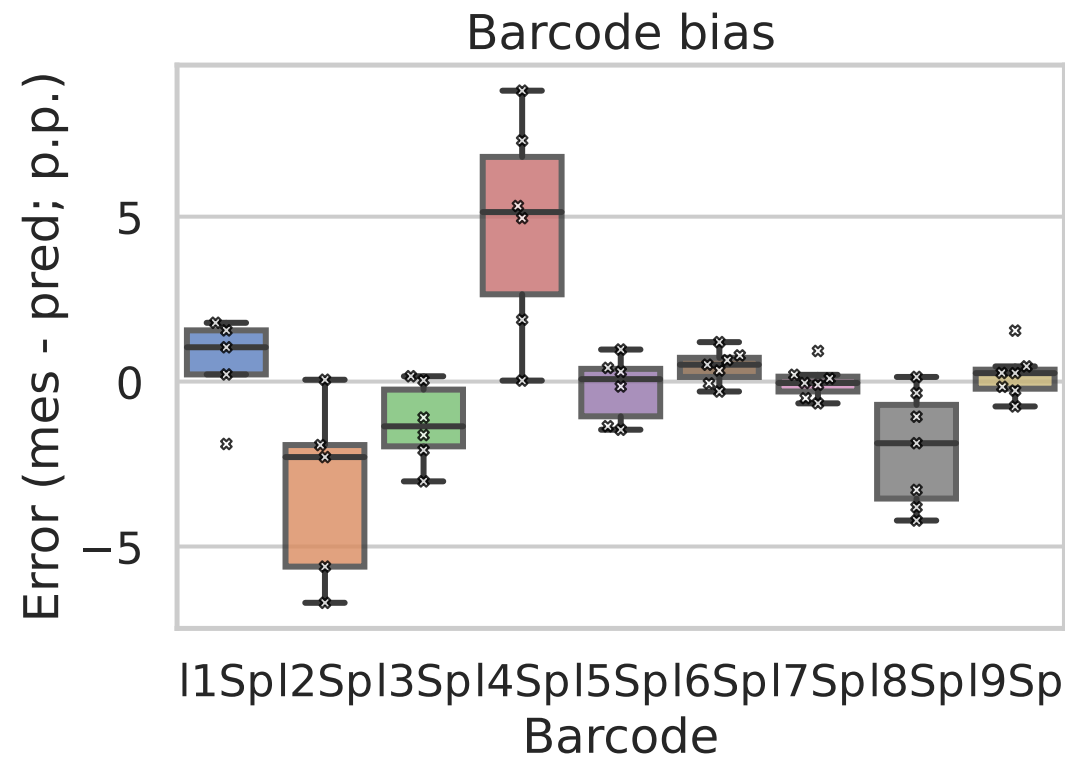
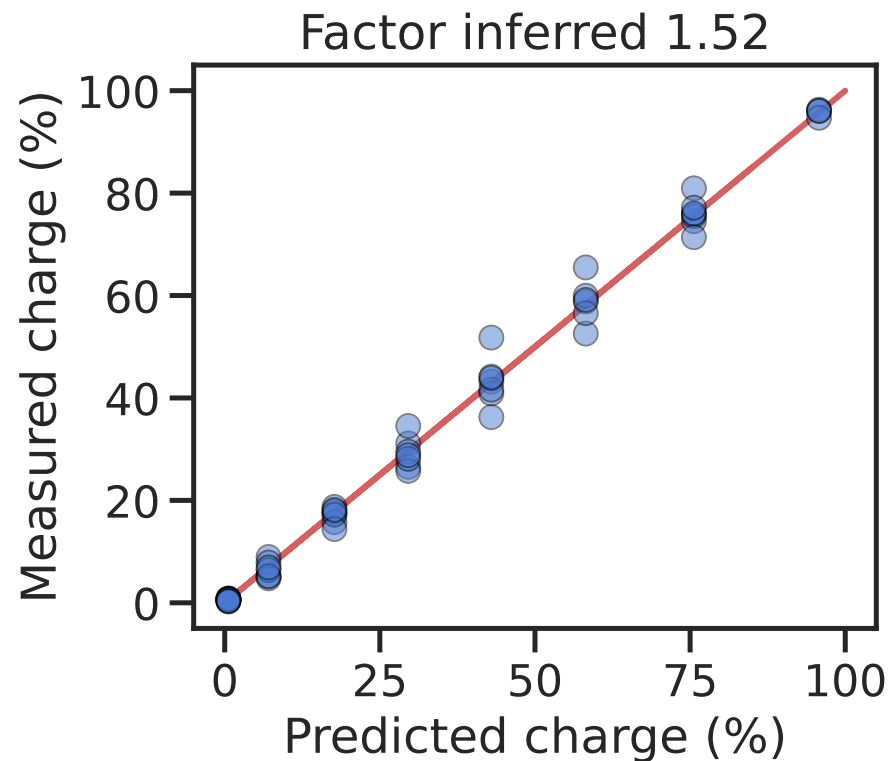
His-GTG-1-1



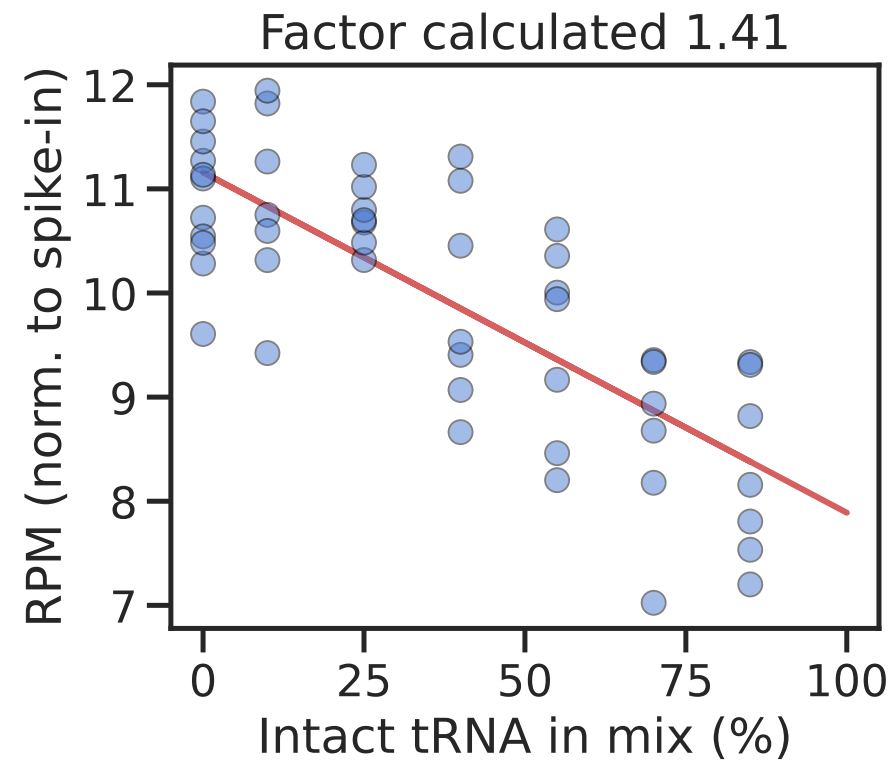
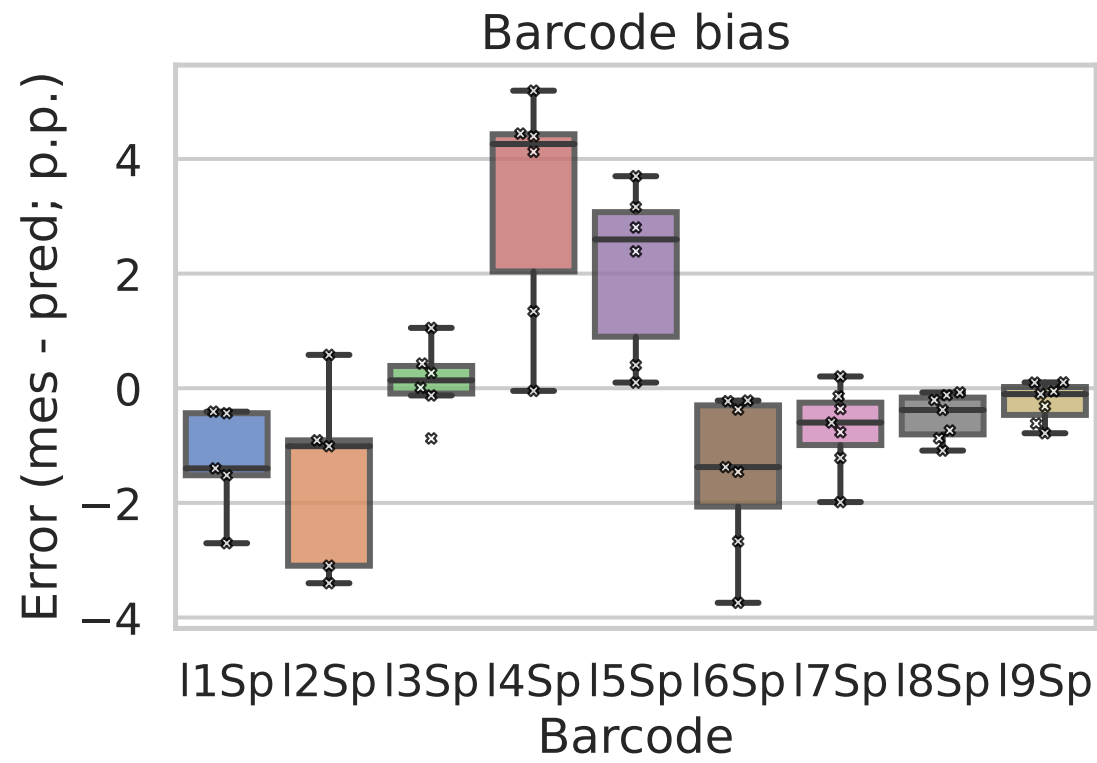
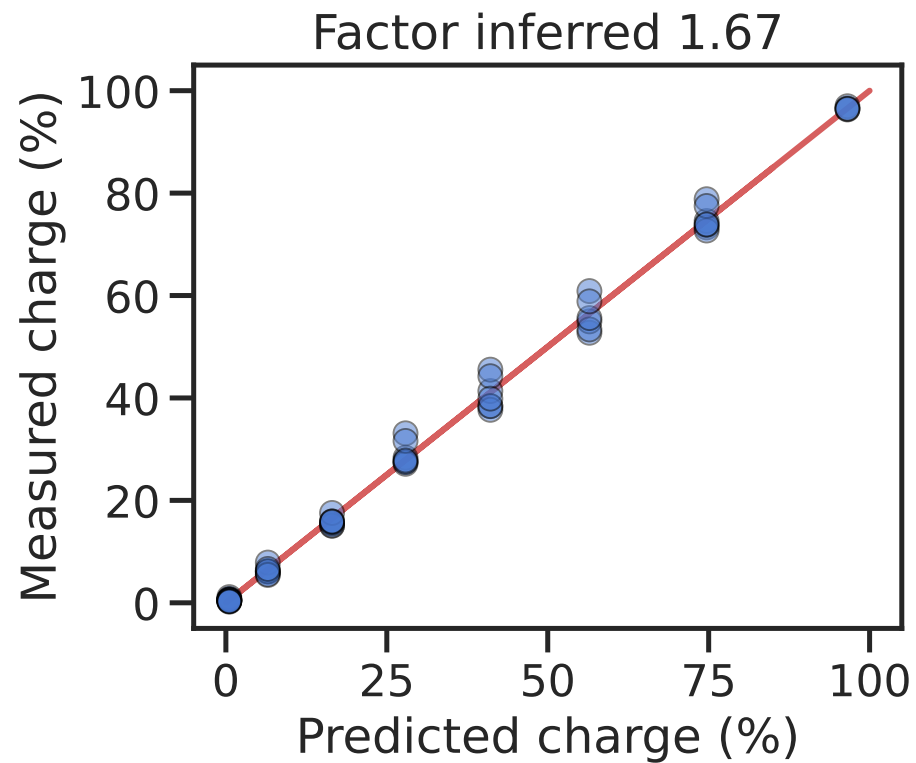
Ile-AAT-7-1



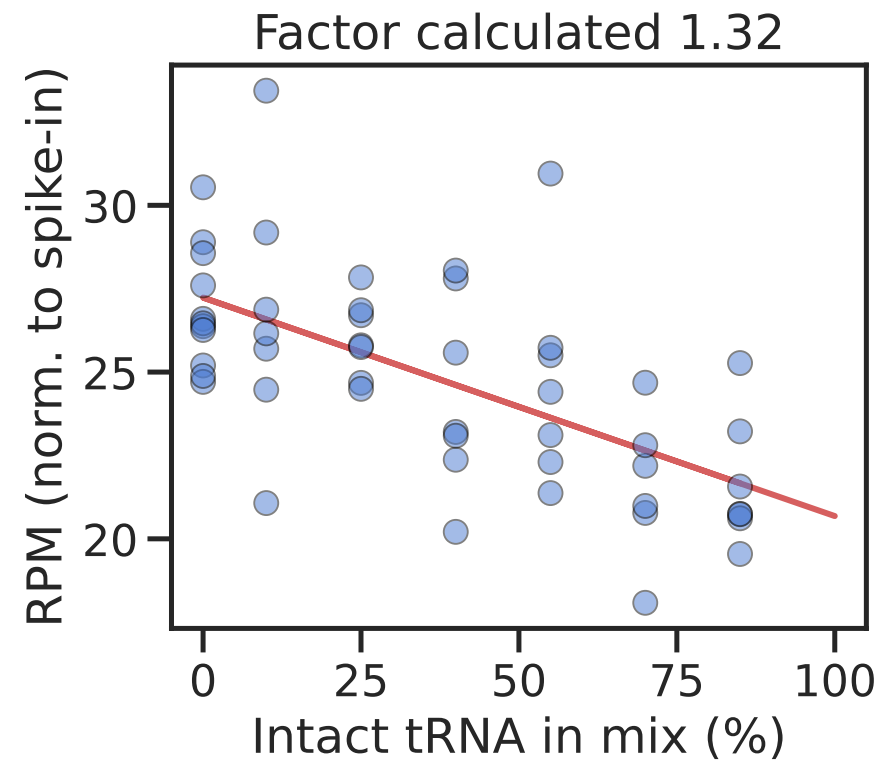
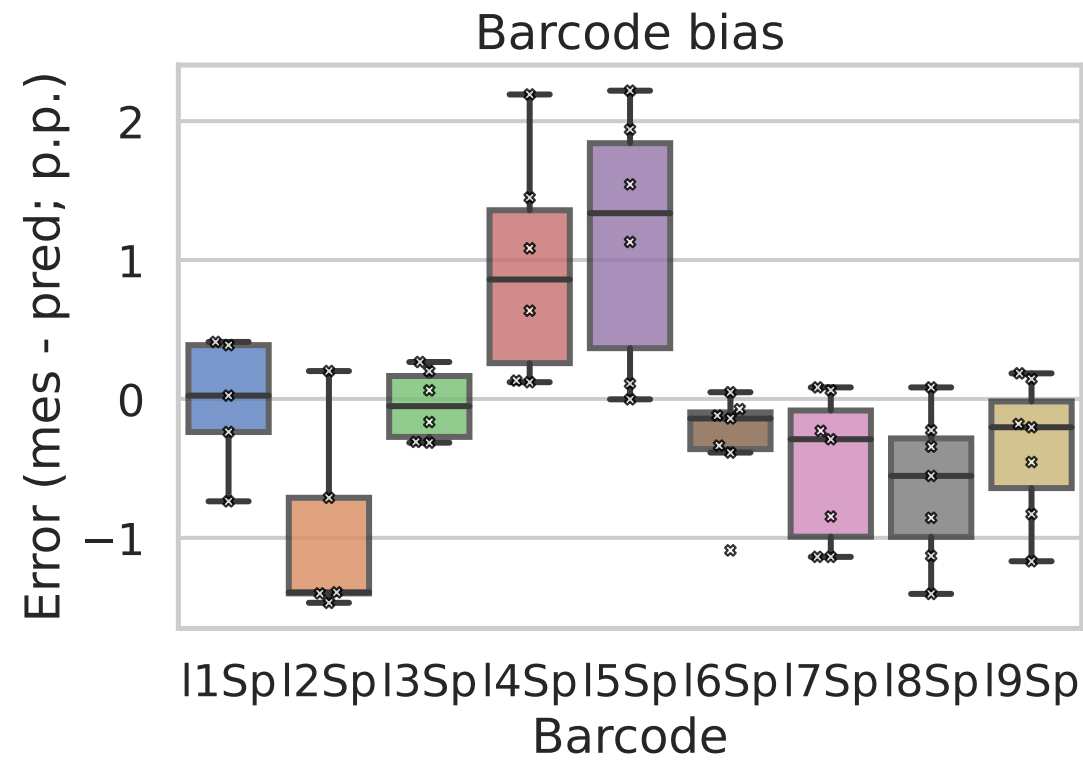
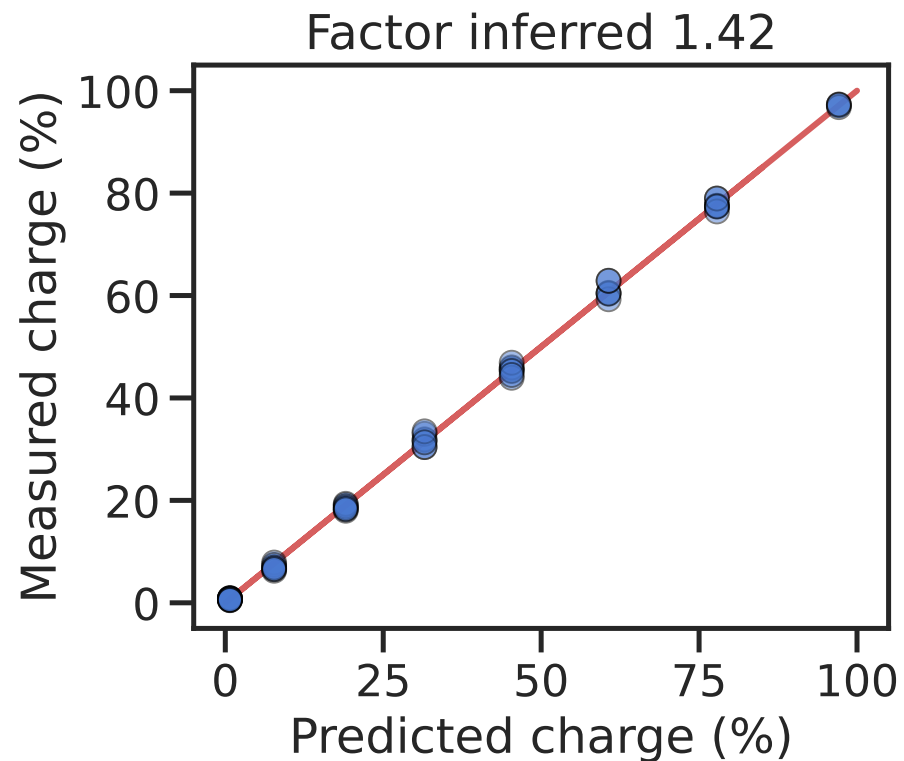
Thr-CGT-2-1



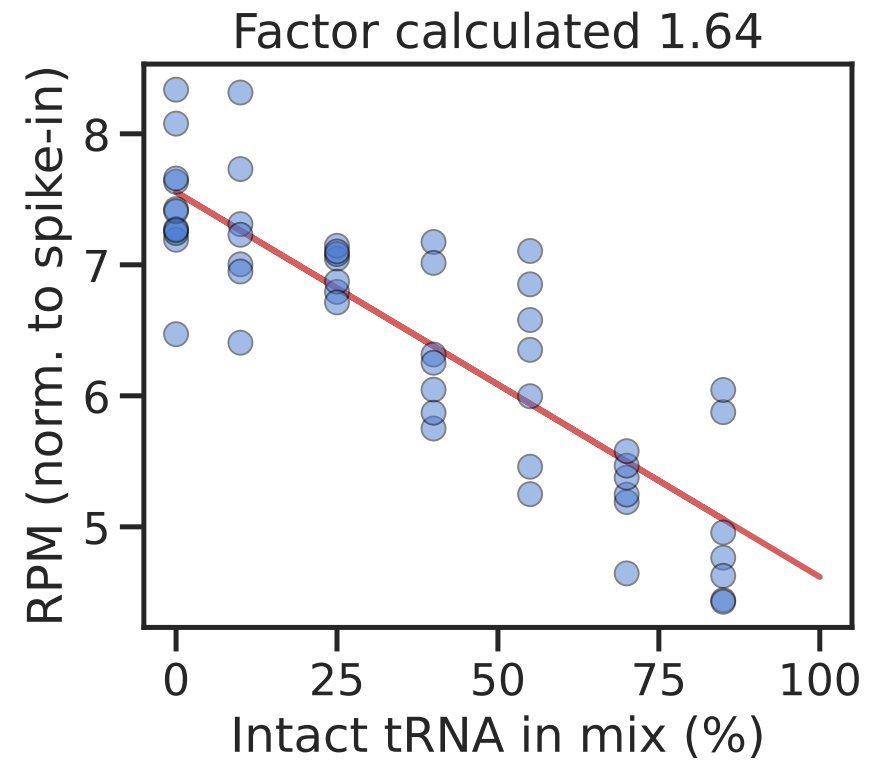
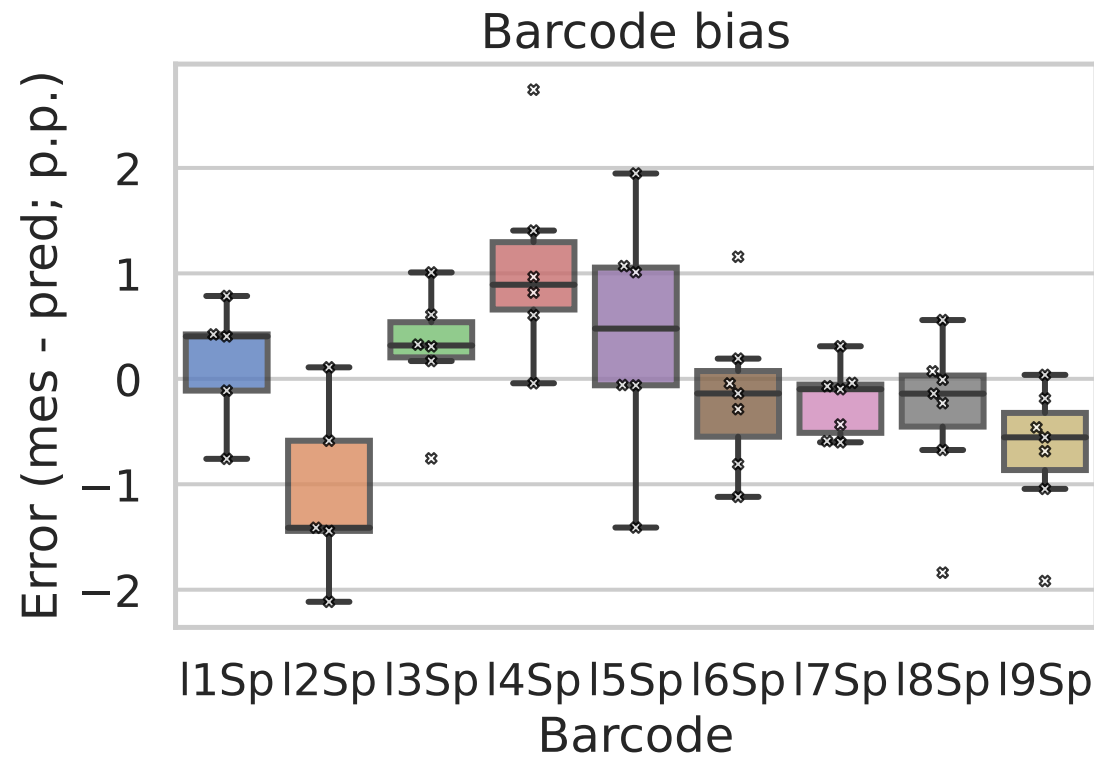
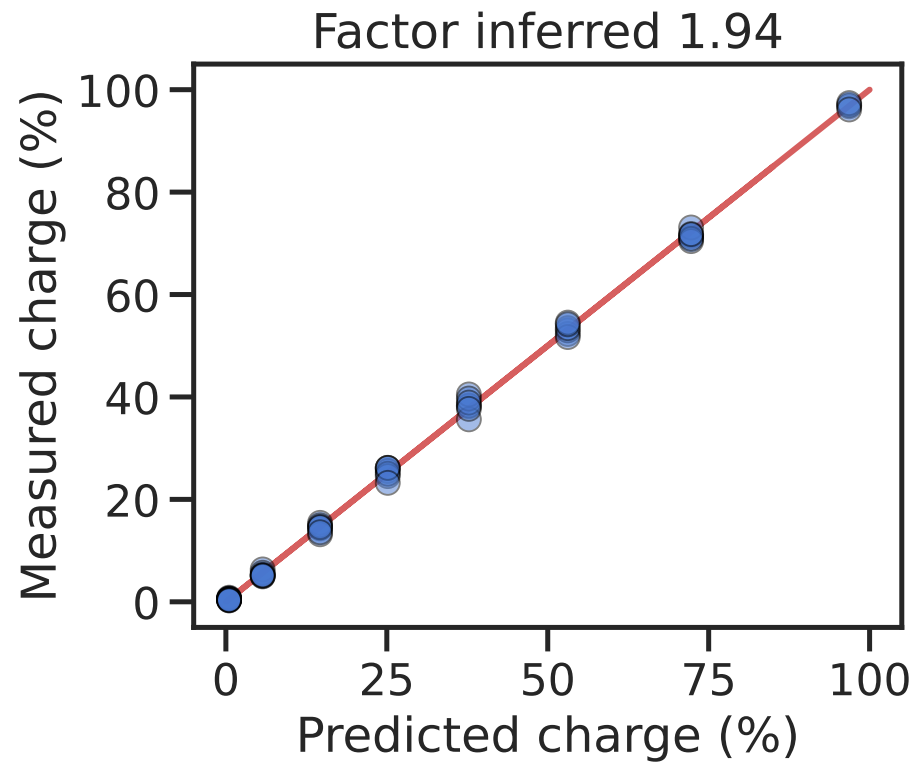
Arg-TCG-1-1



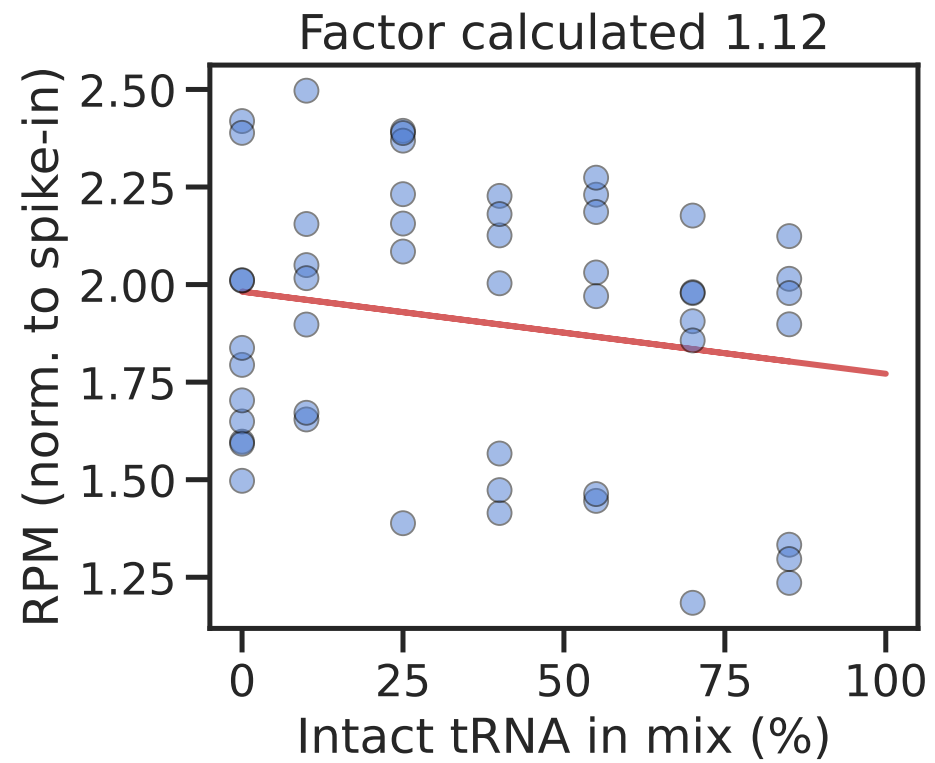
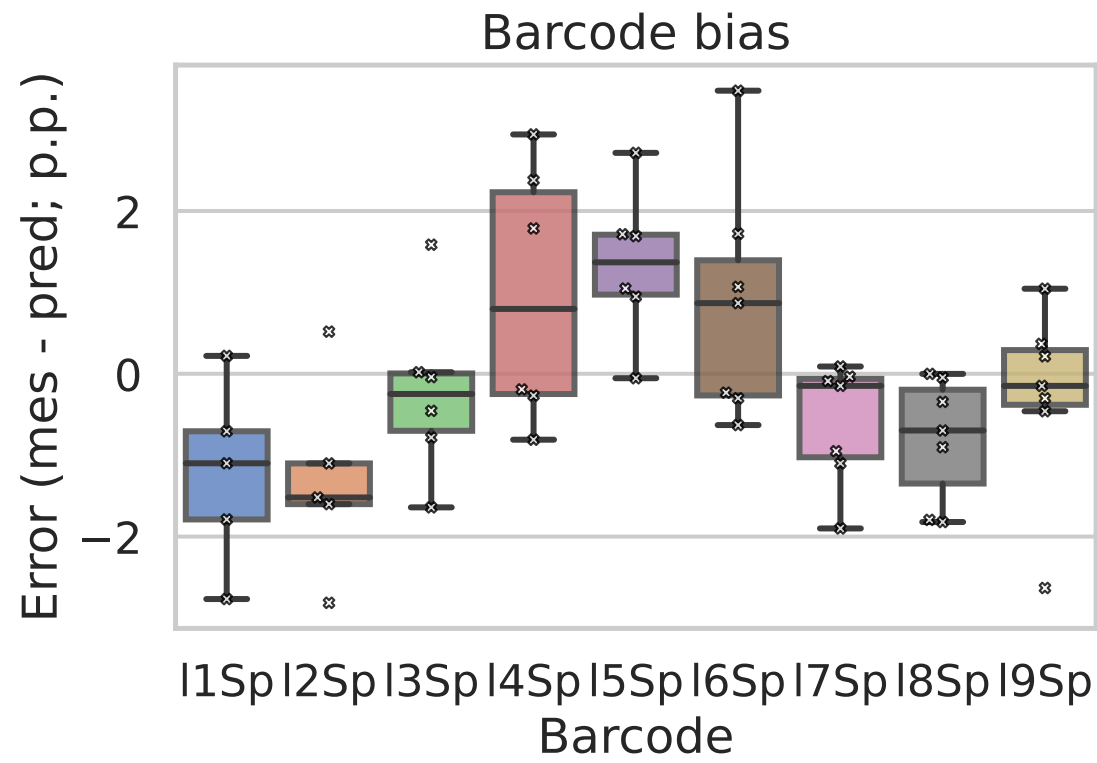
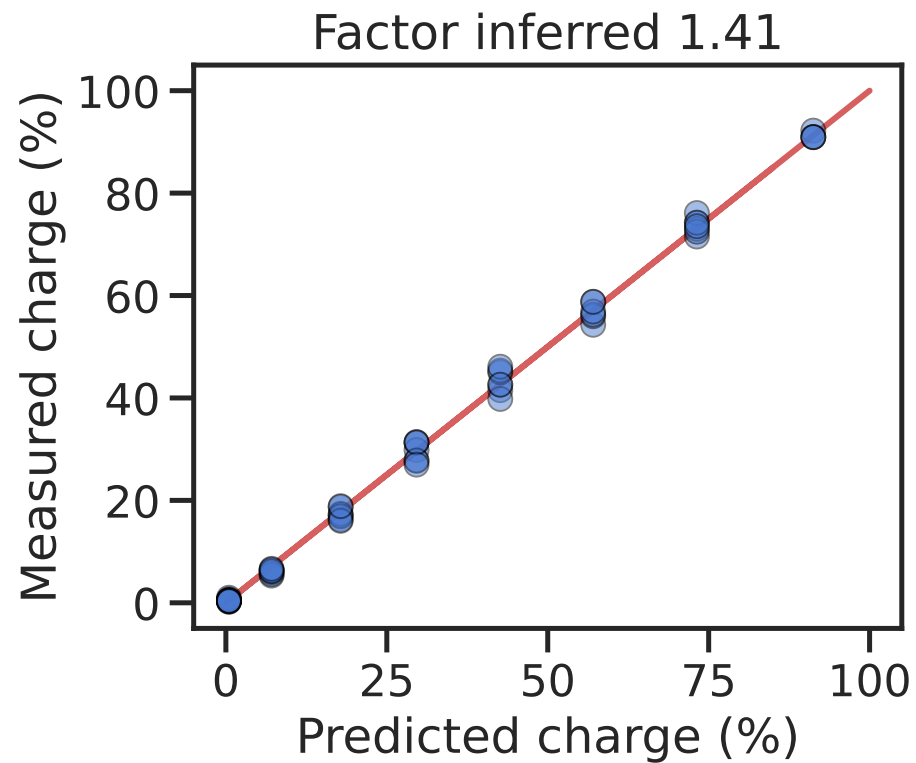
Lys-CTT-1-1



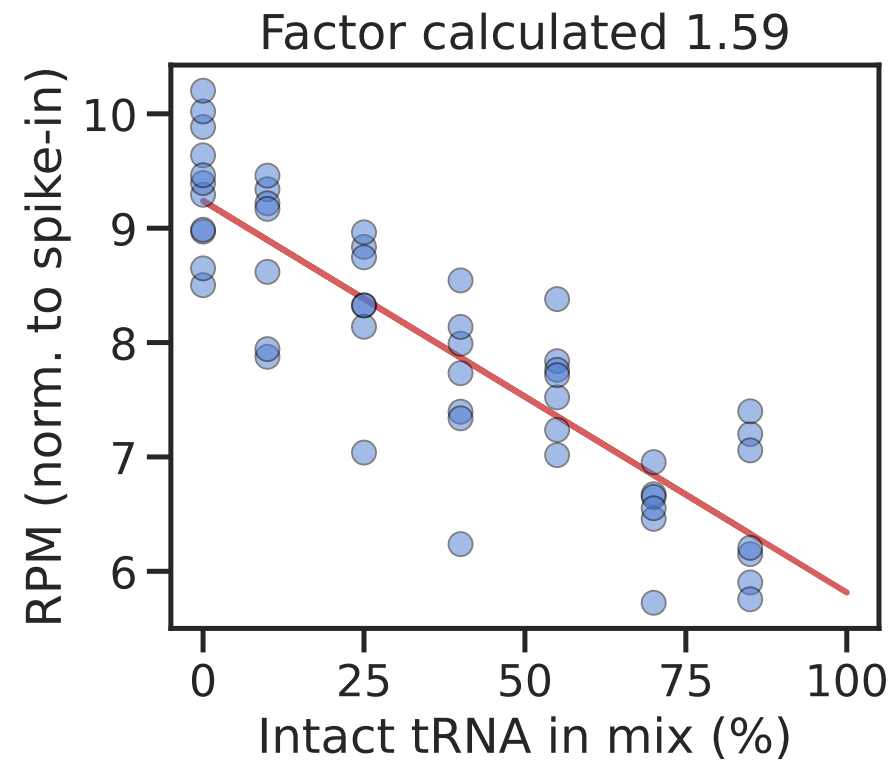
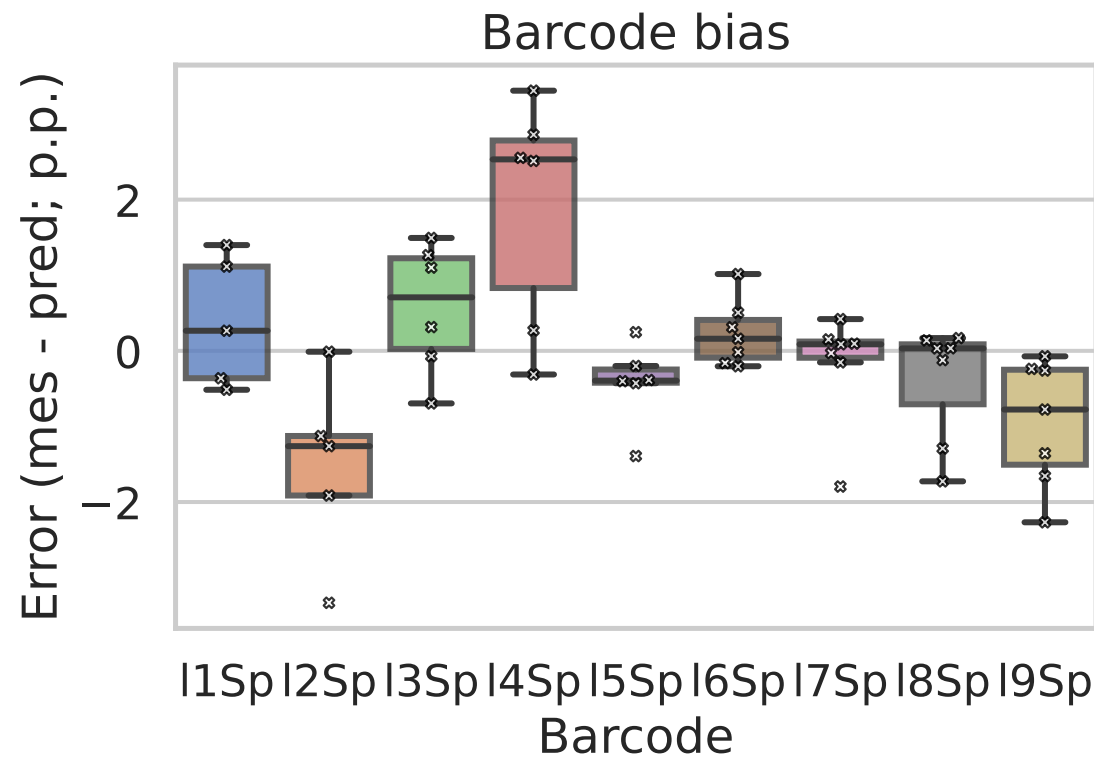
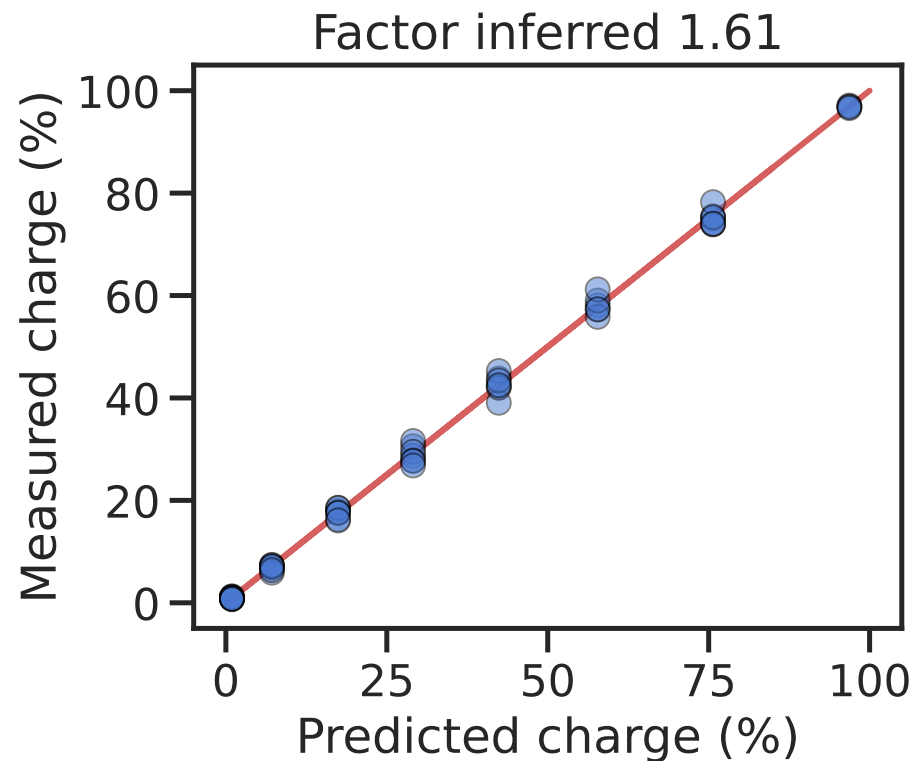
Leu-CAA-4-1



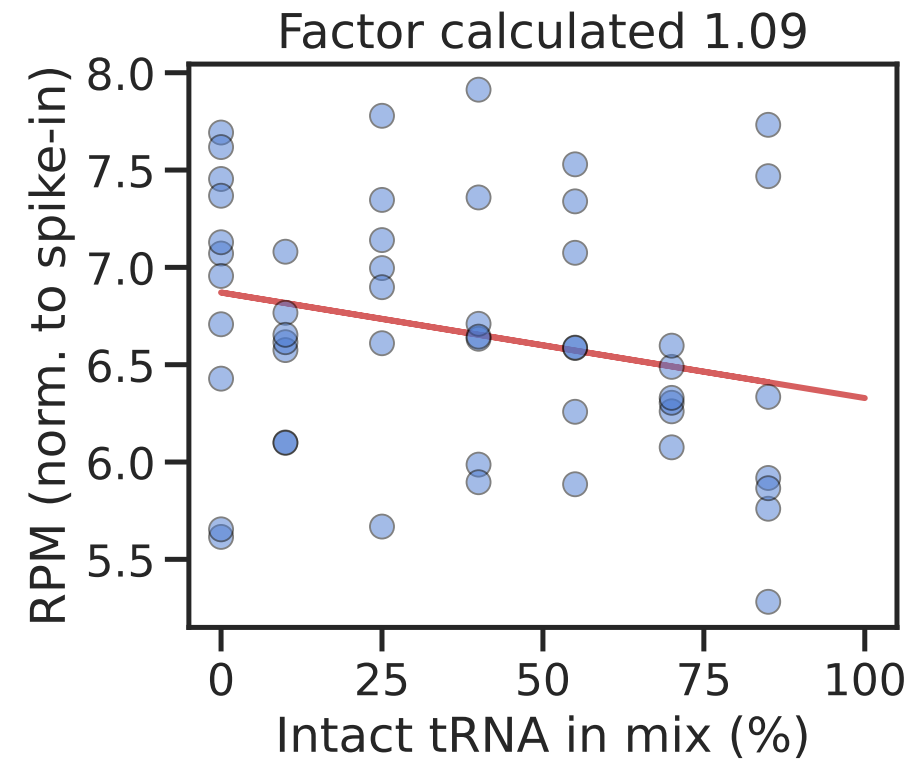
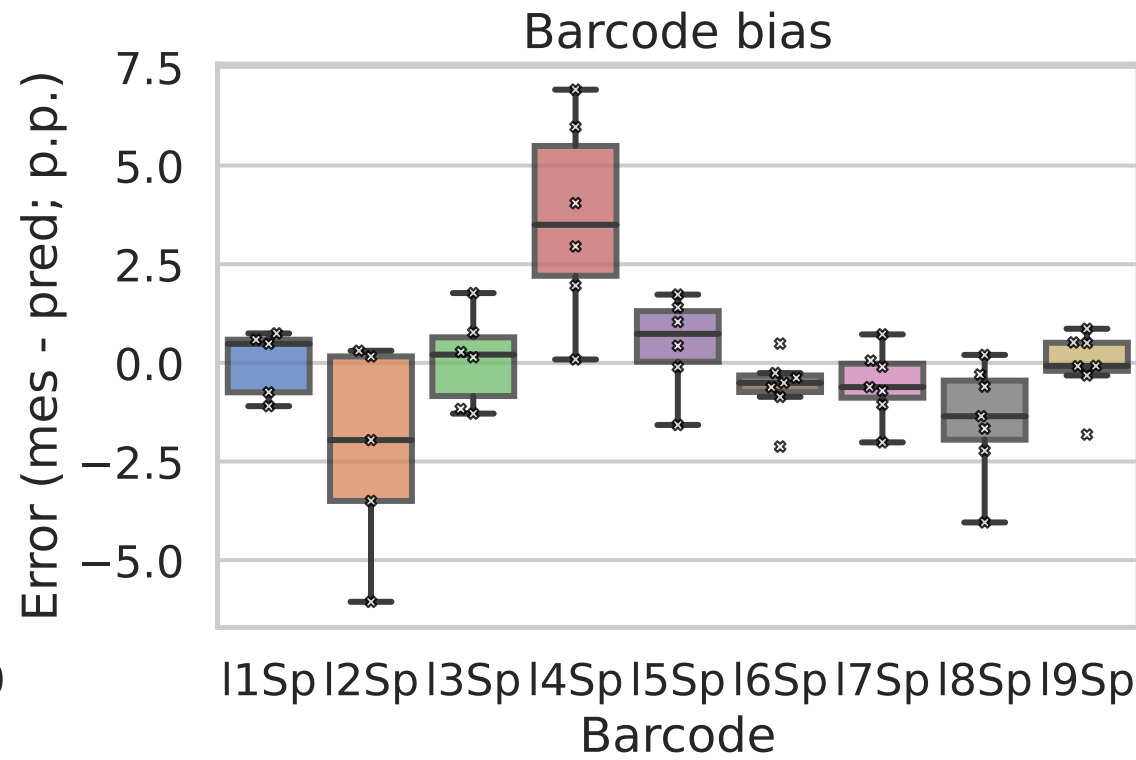
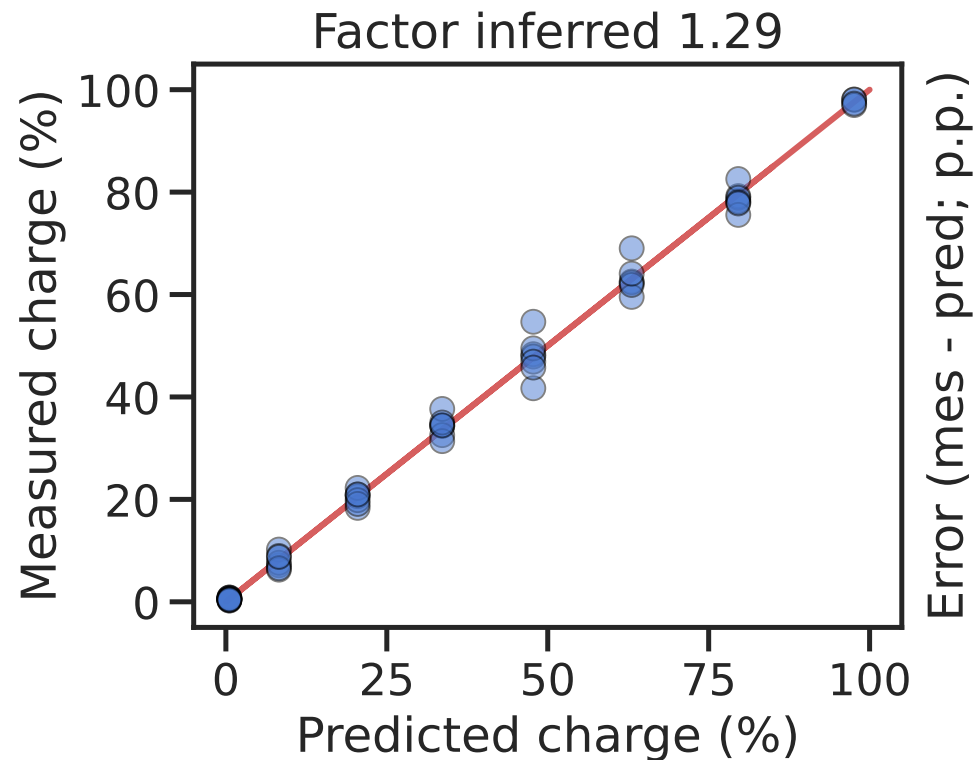
SeC-TCA-1-1



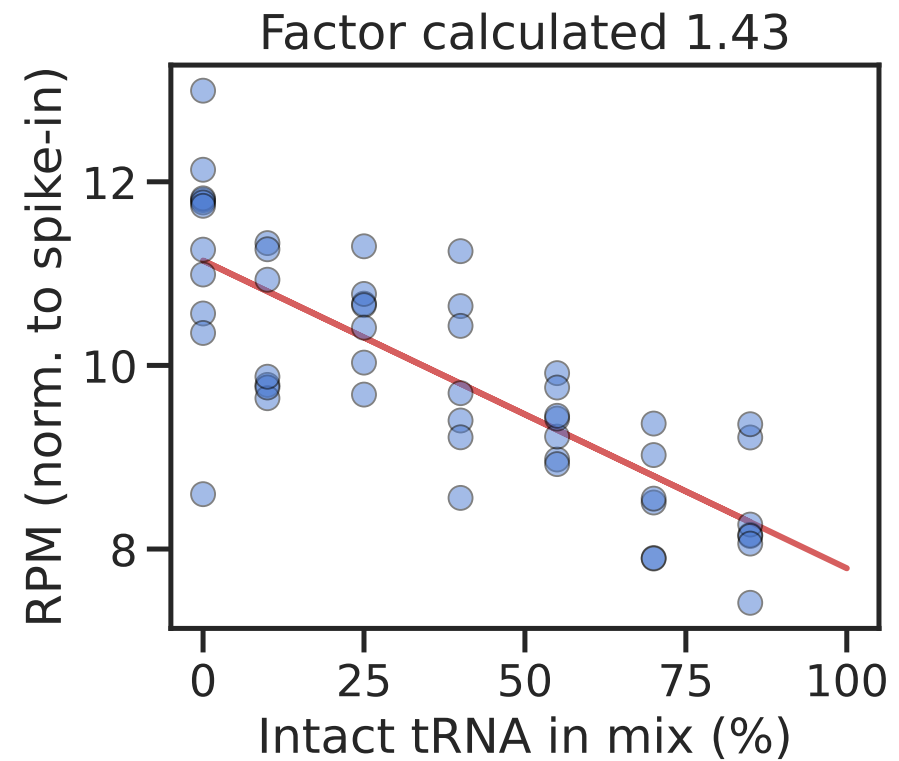
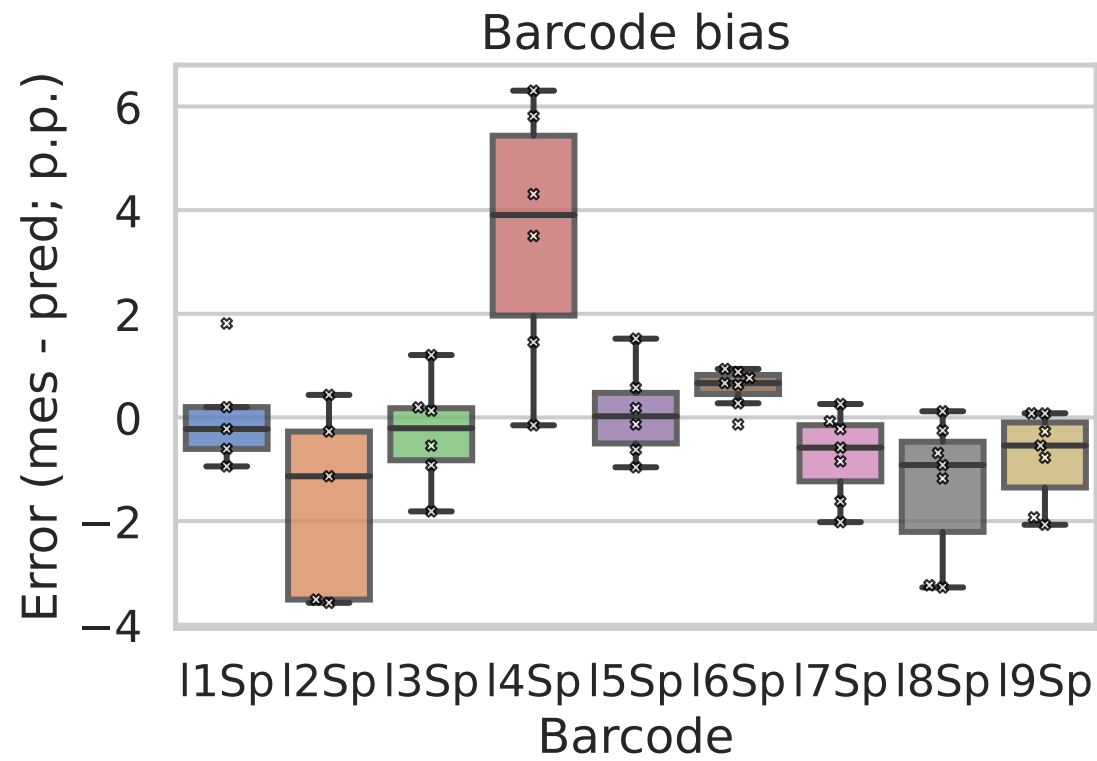
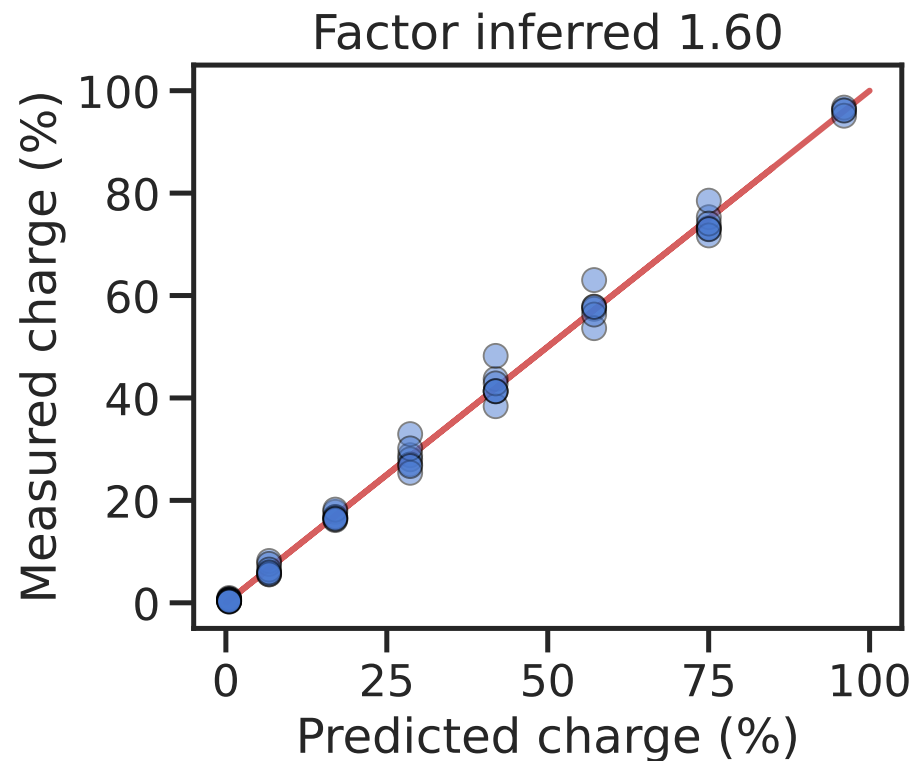
Val-TAC-2-1



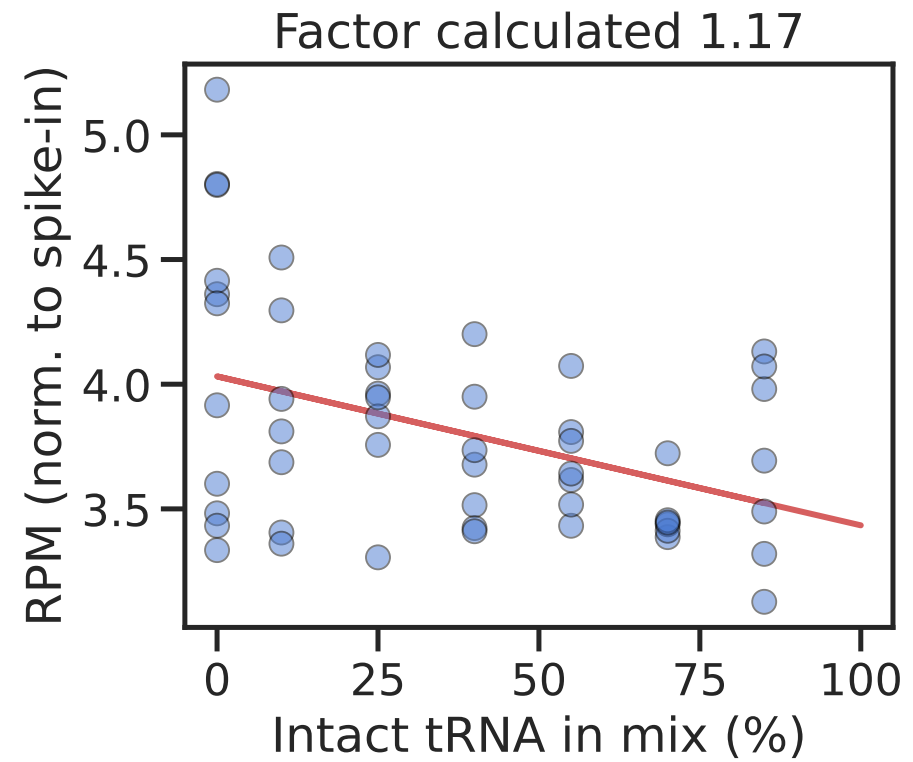
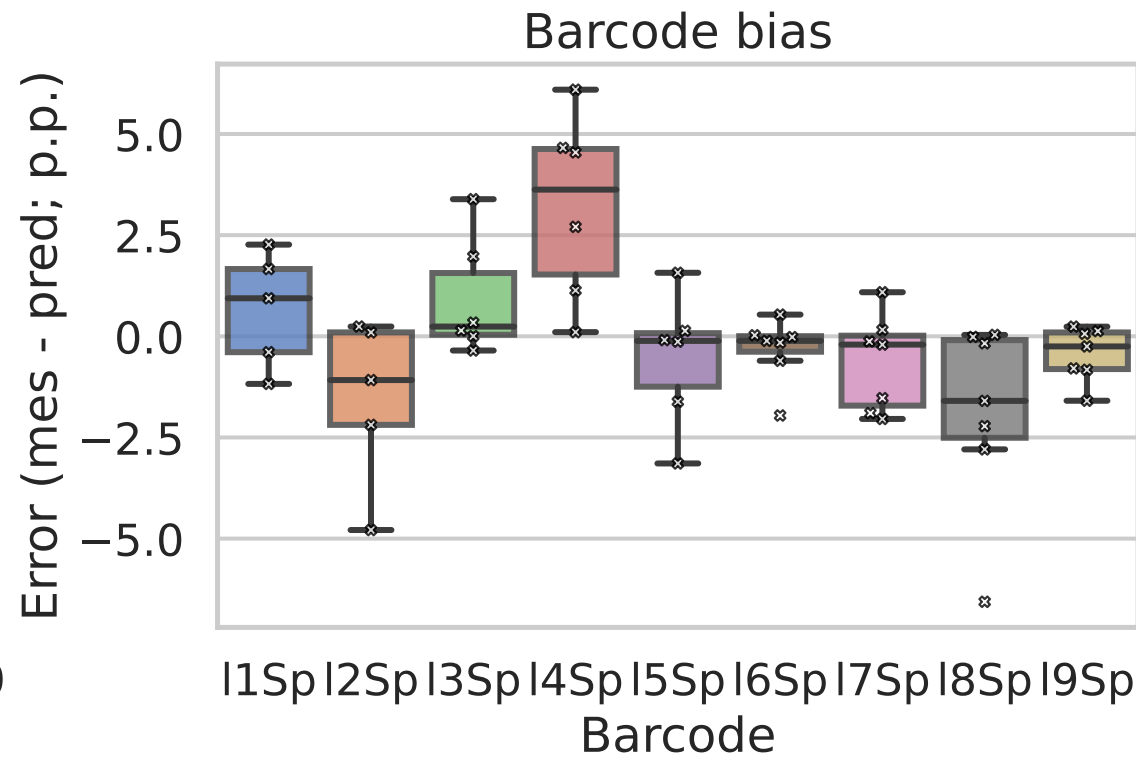
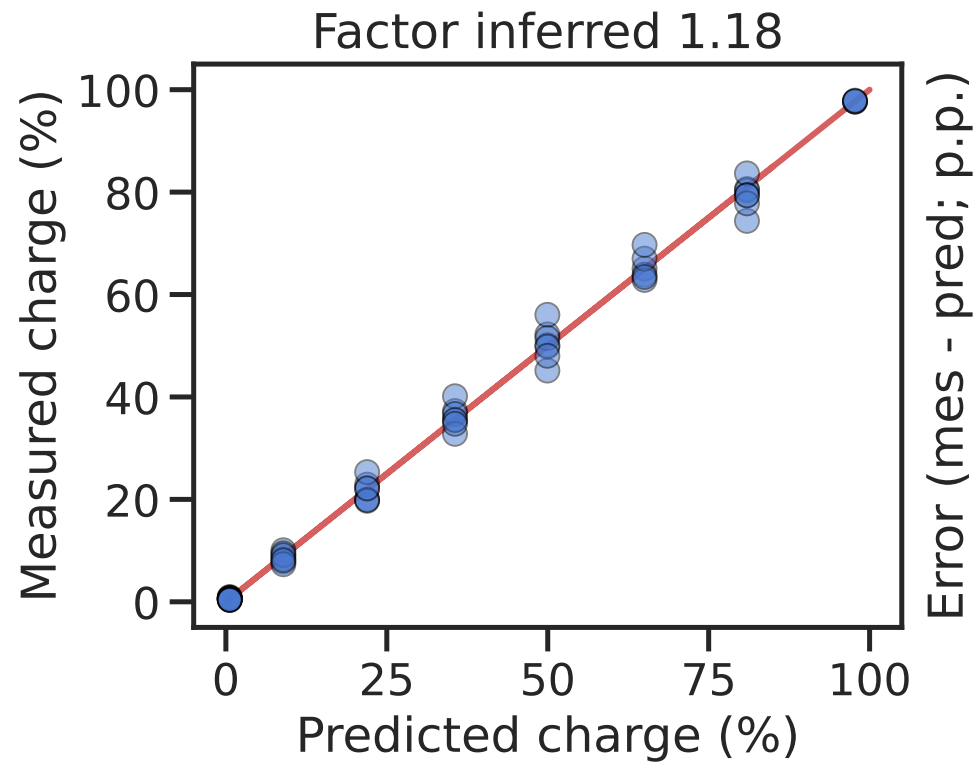
Gln-TTG-2-1



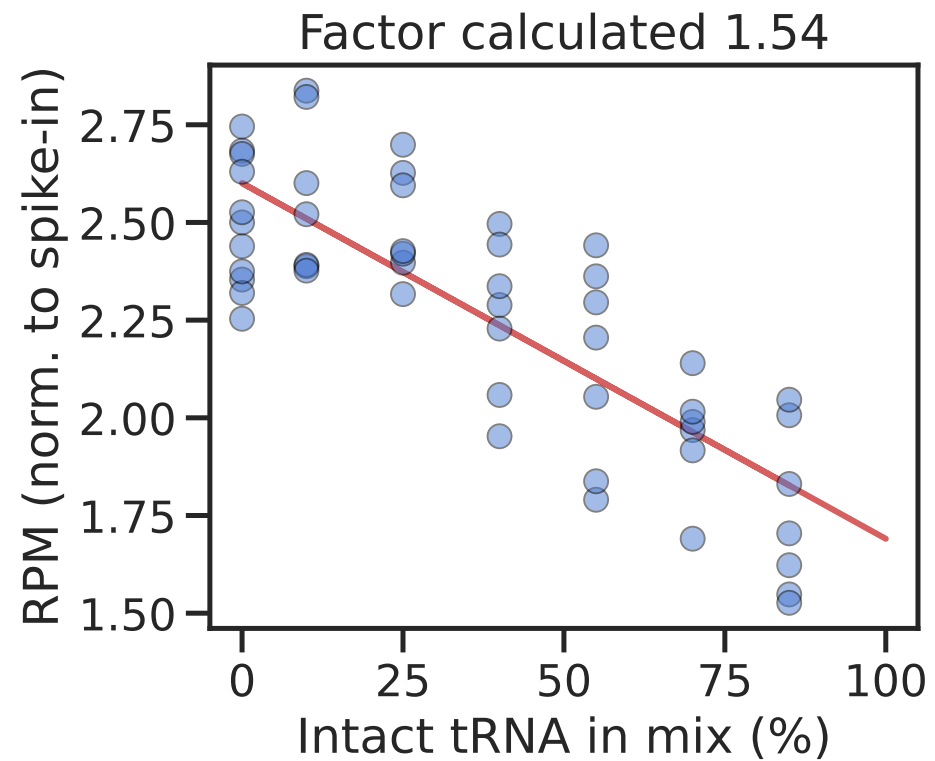
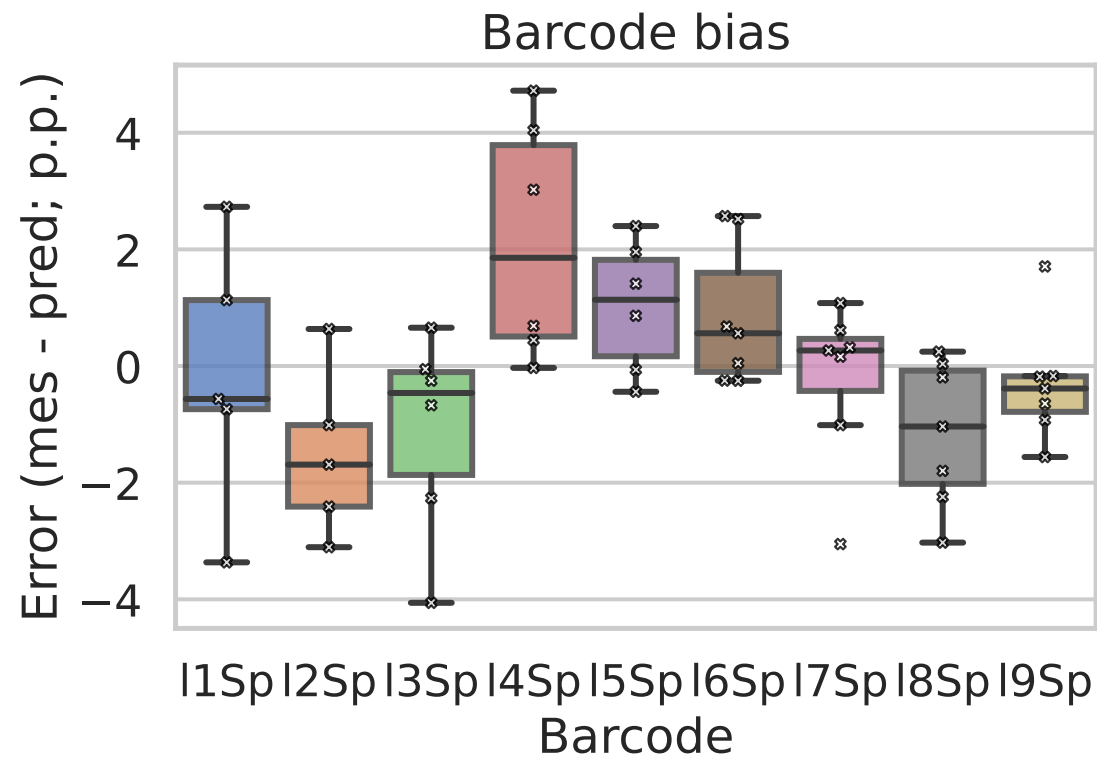
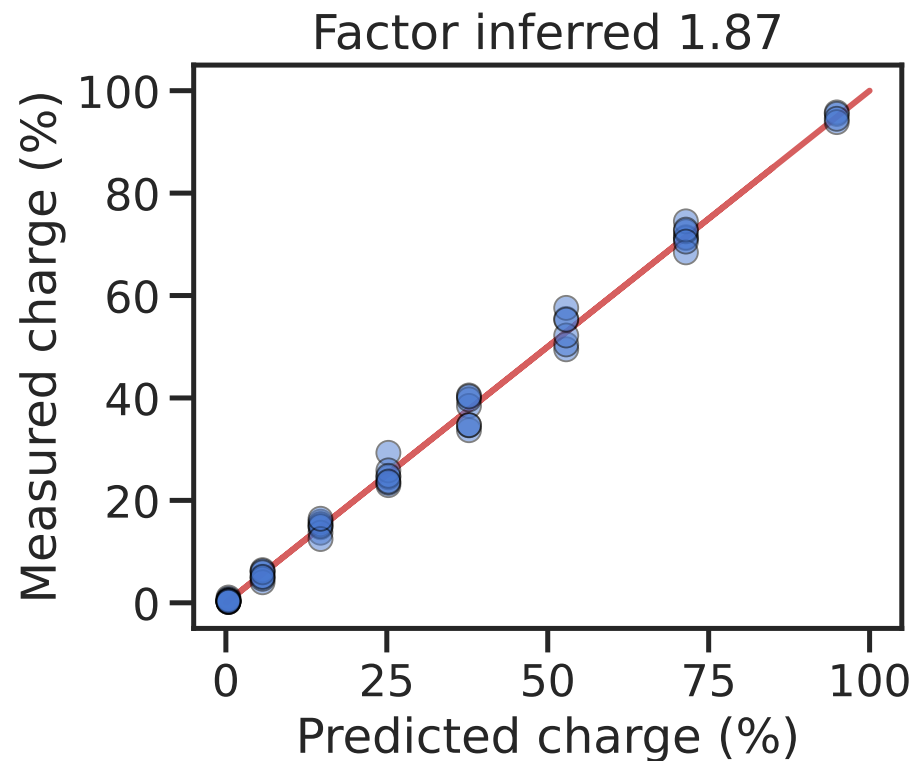
Gln-CTG-2-1



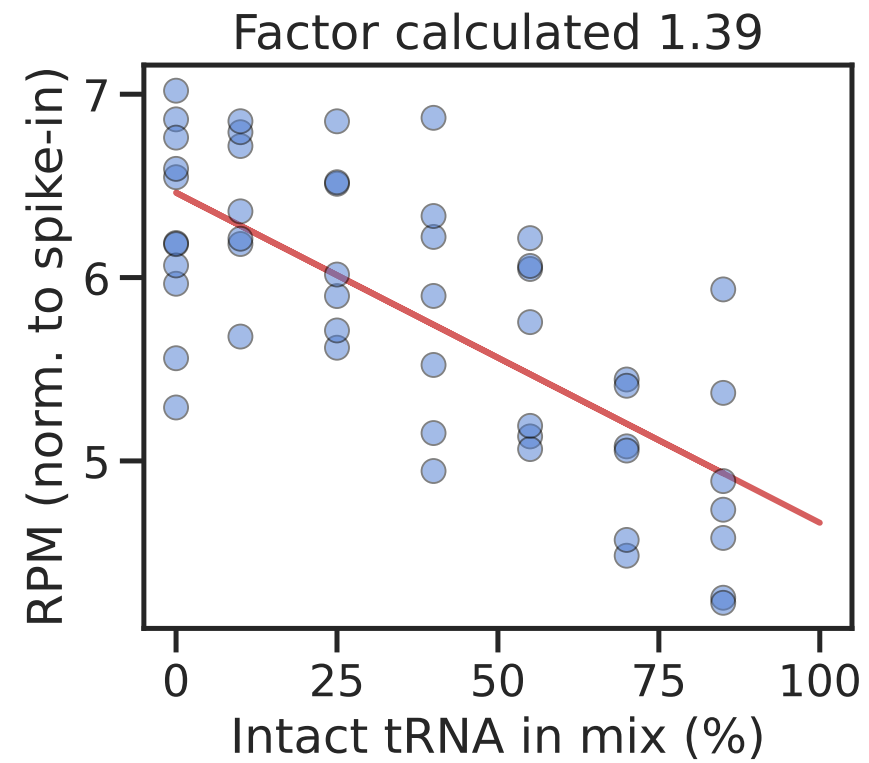
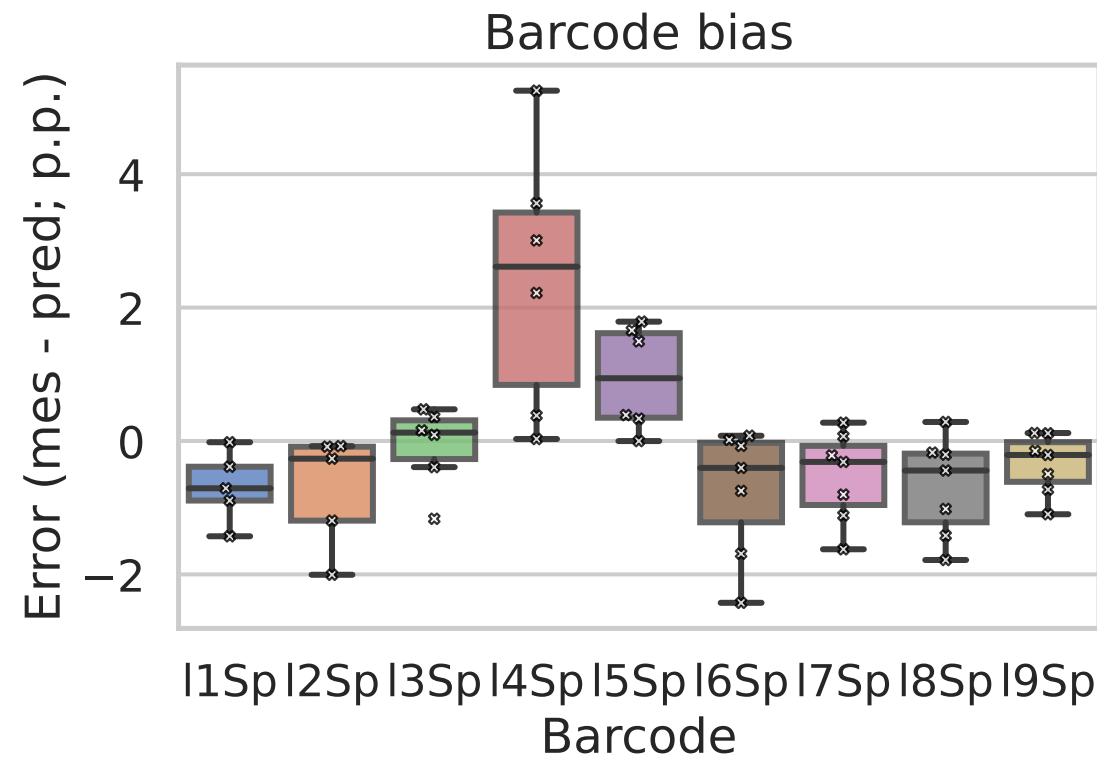
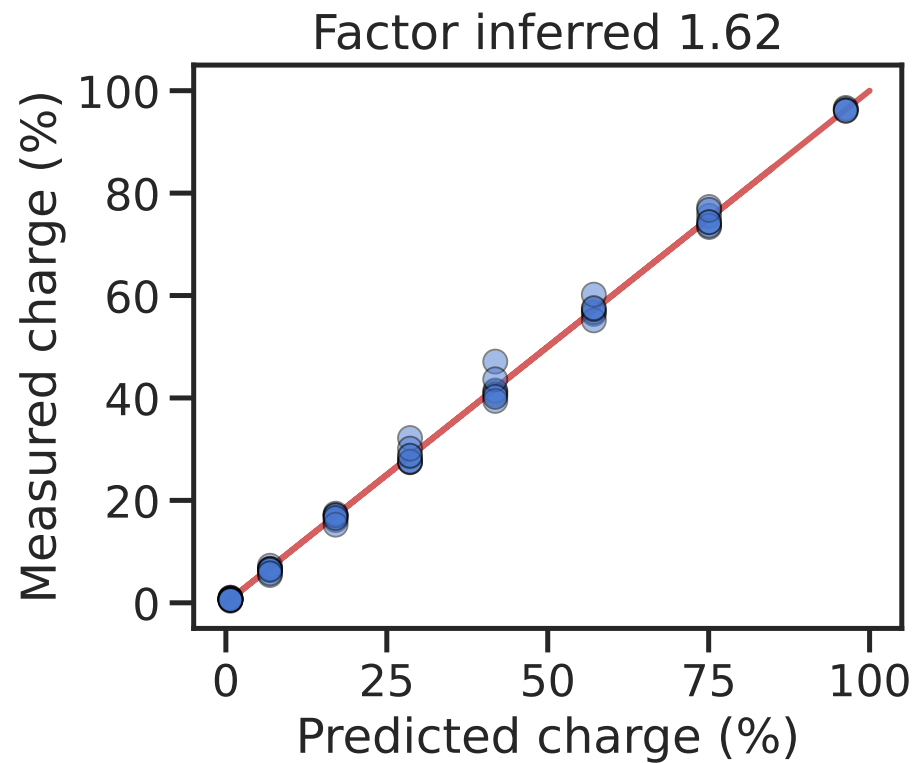
Gln-TTG-1-1



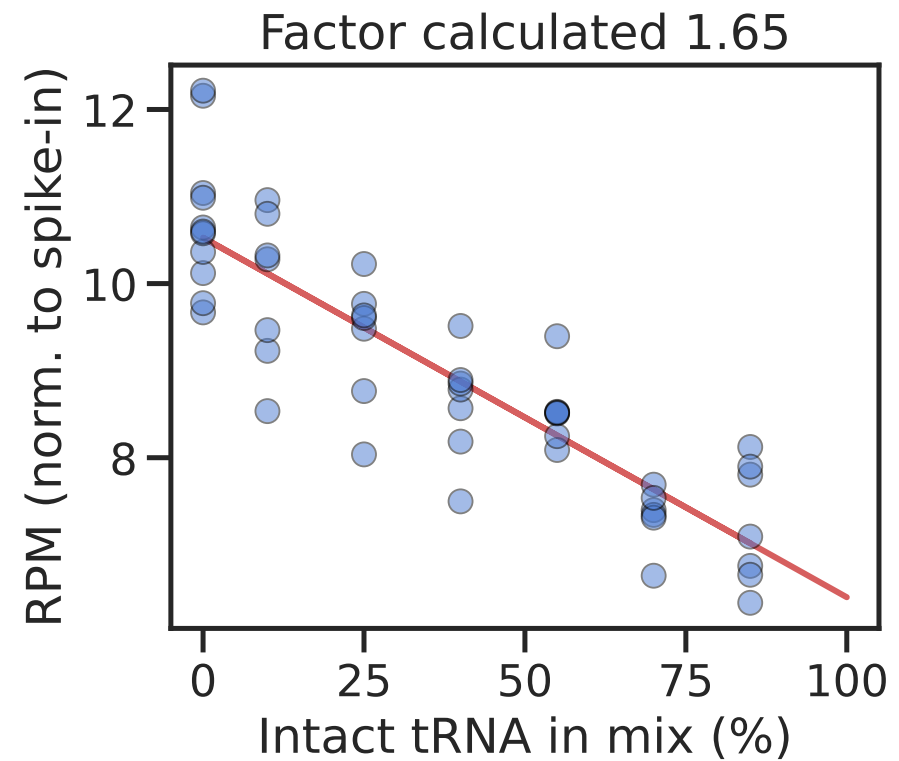
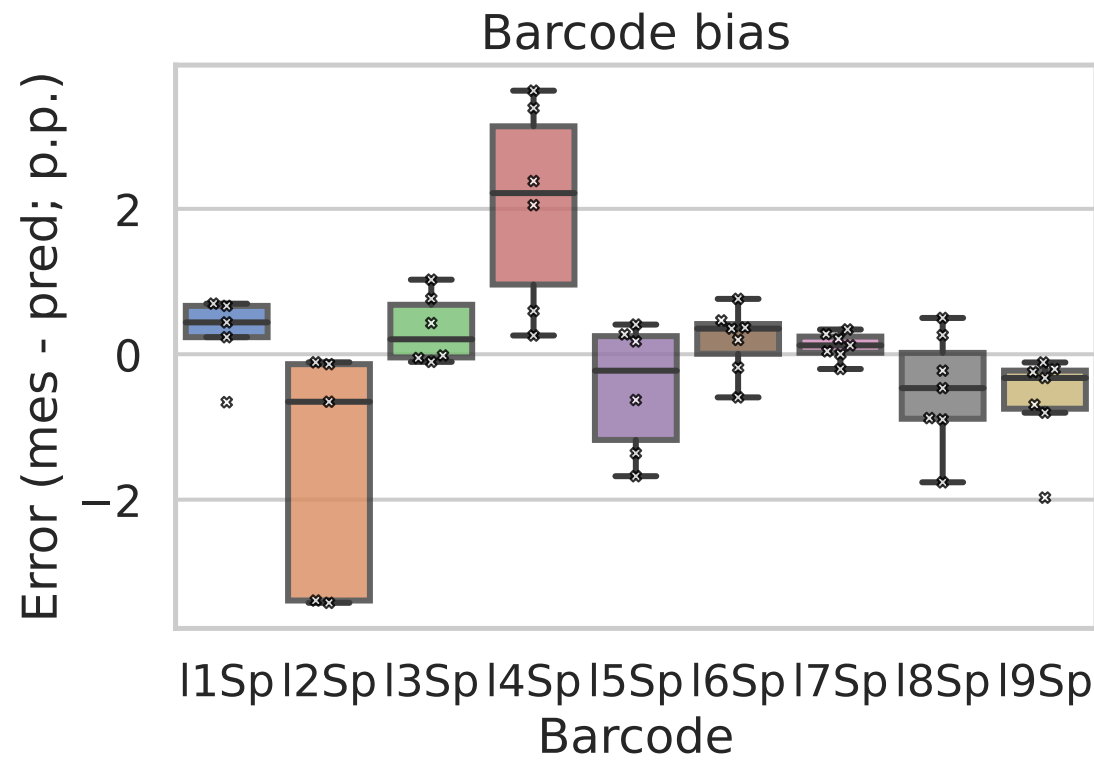
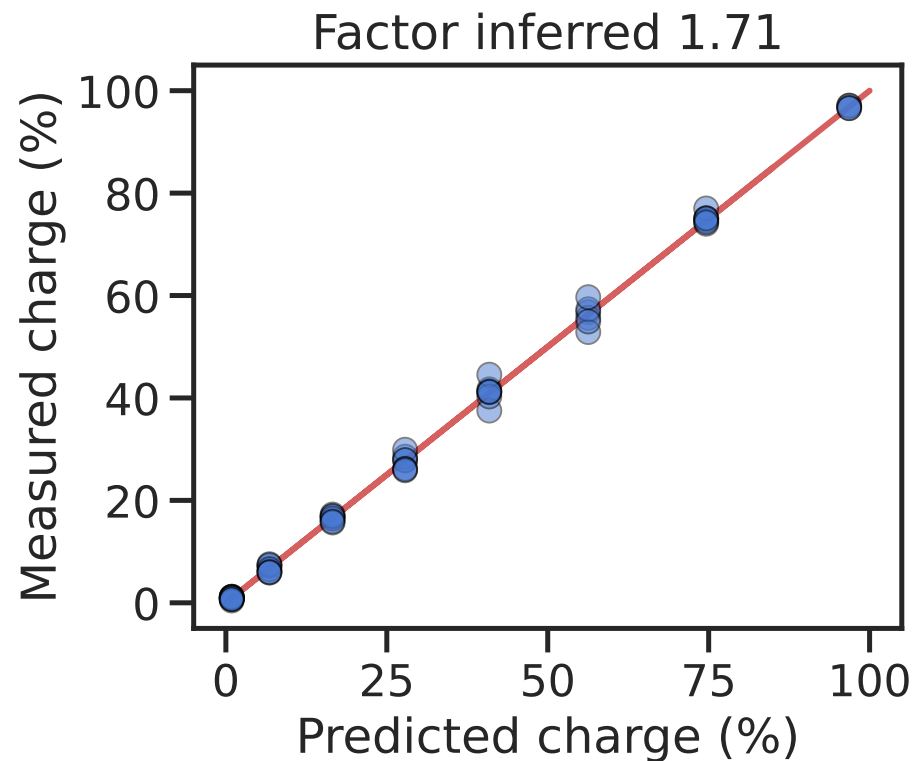
Arg-TCG-4-1



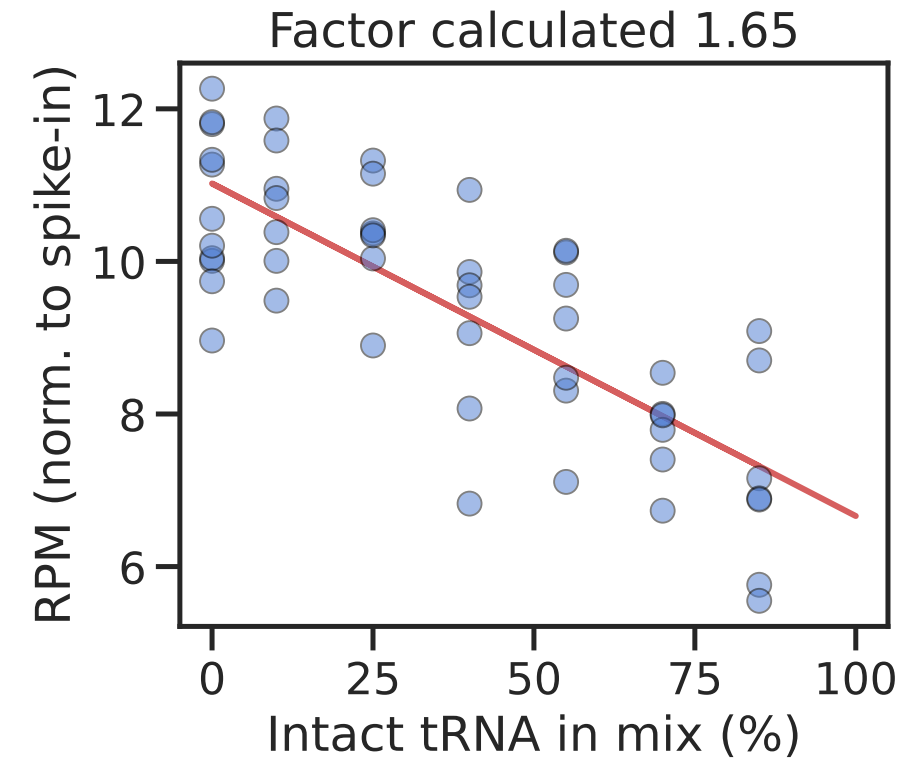
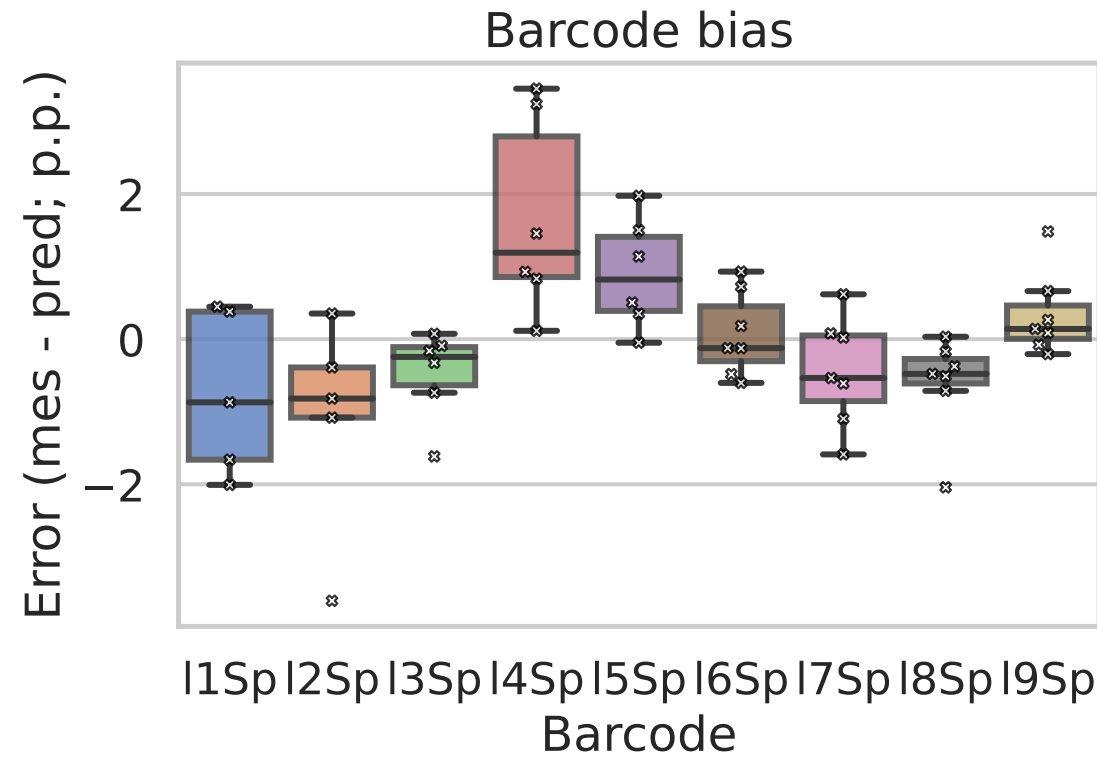
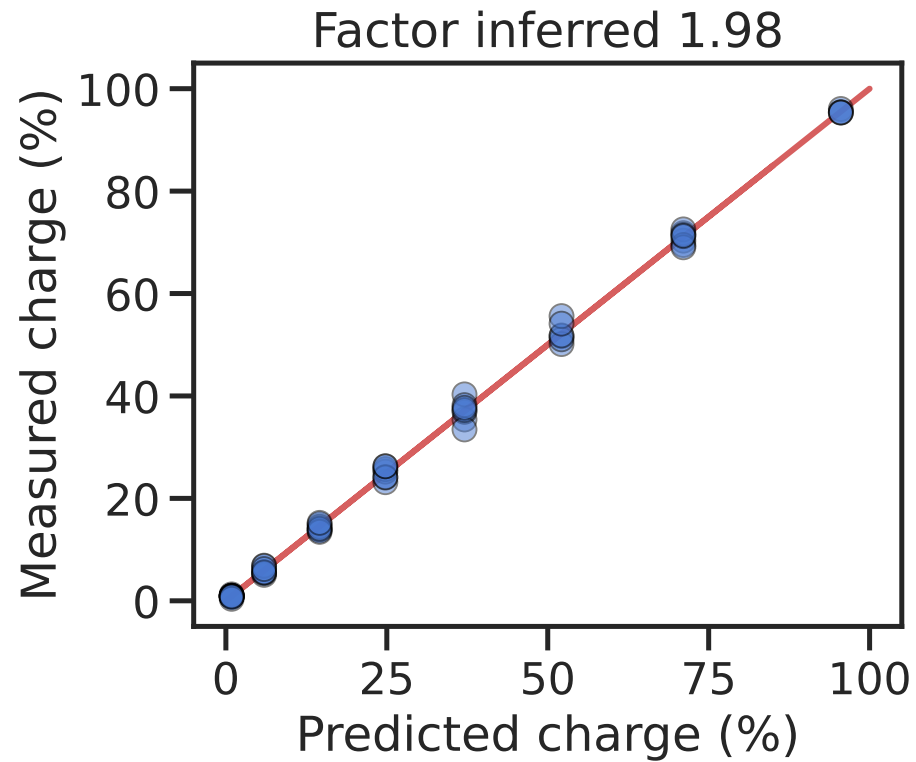
Lys-TTT-6-1



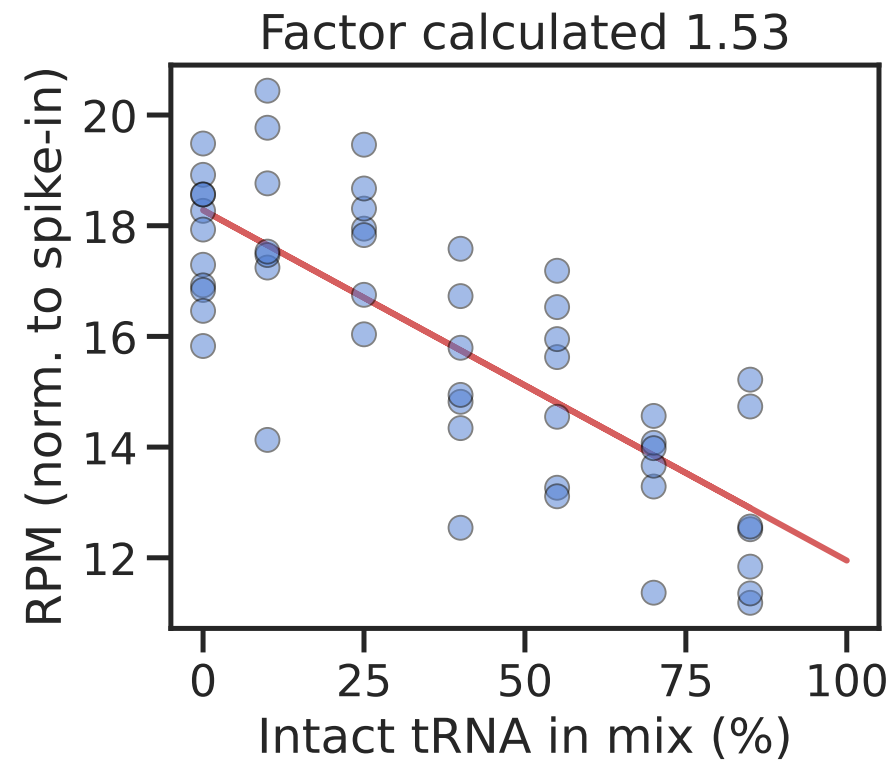
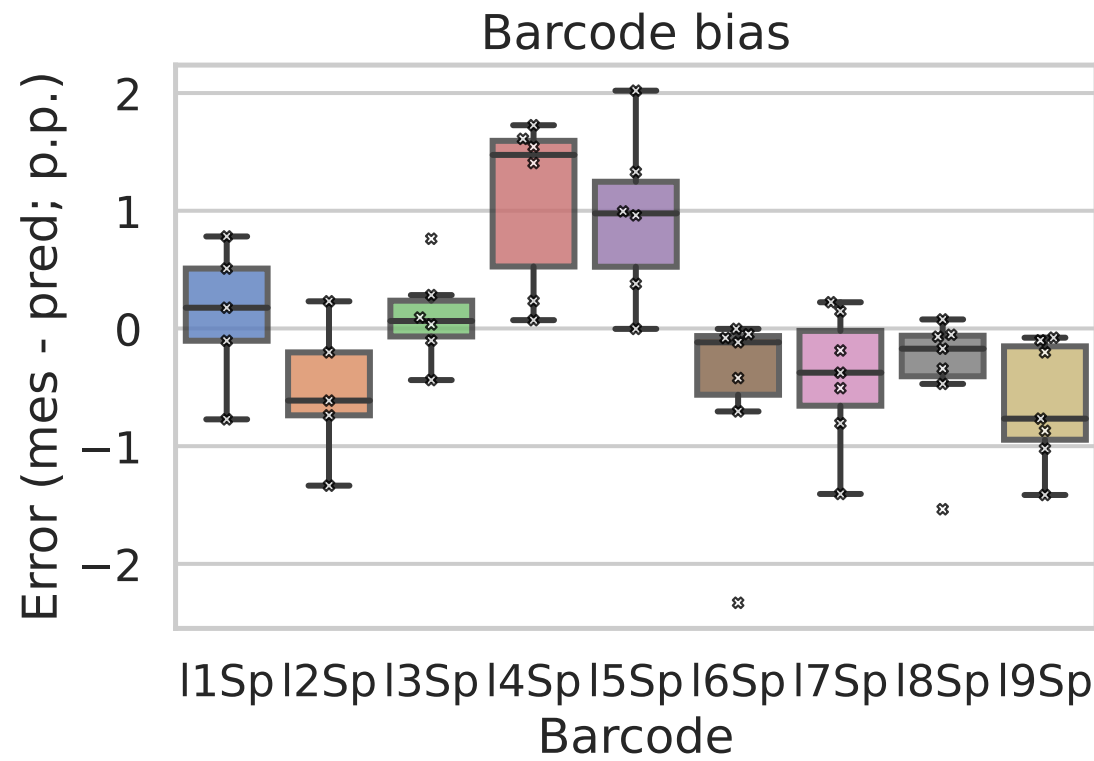
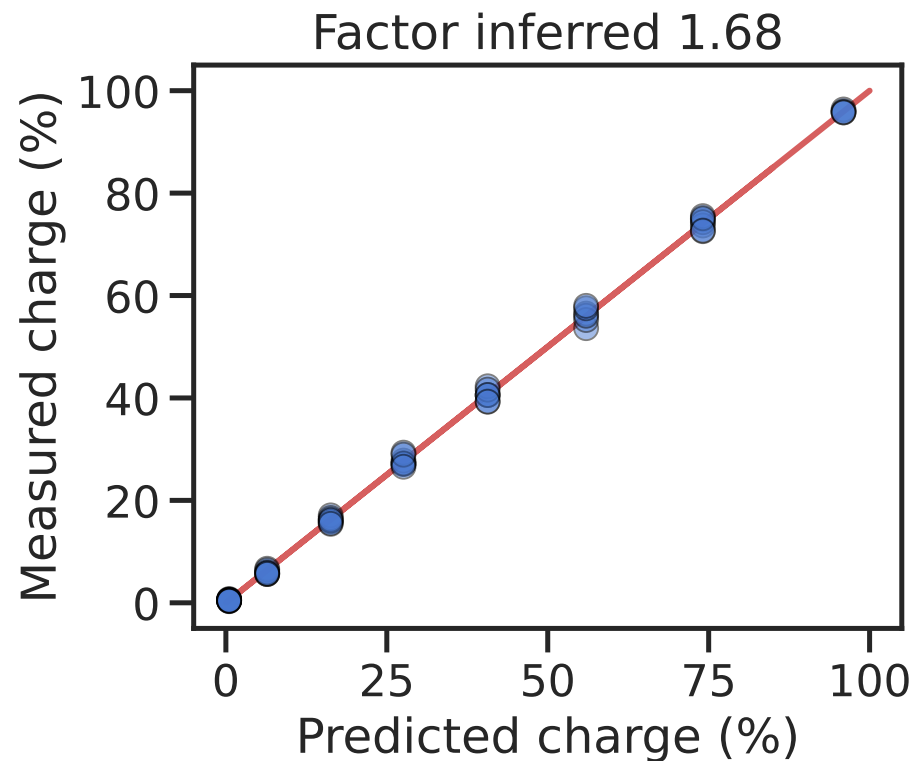
Val-TAC-1-1



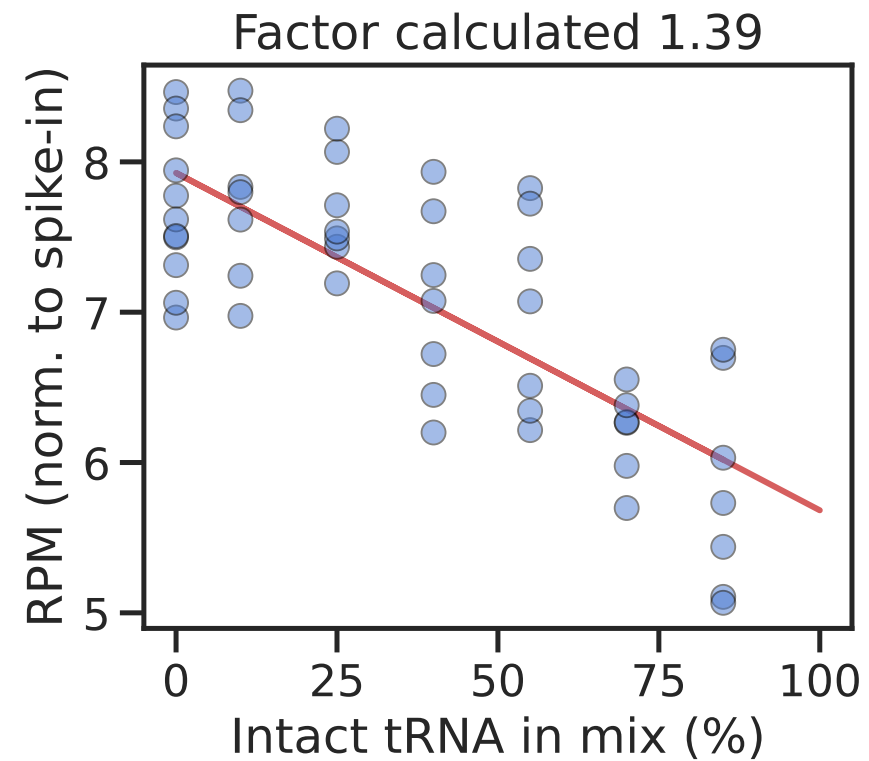
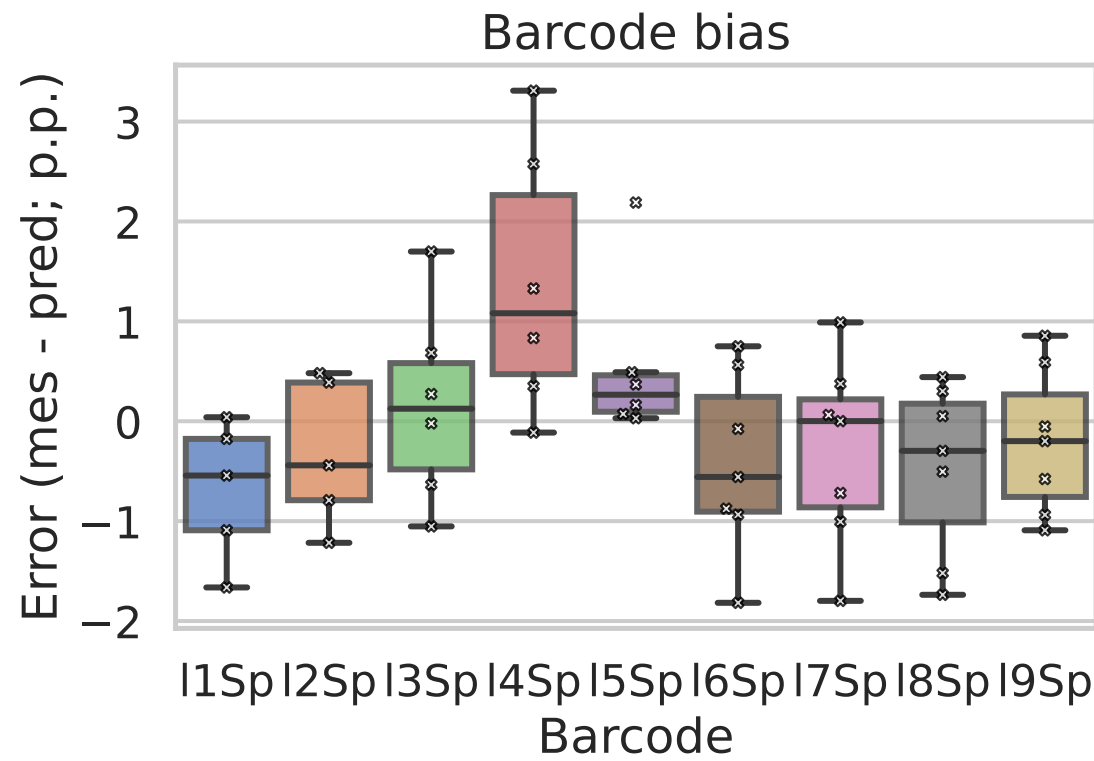
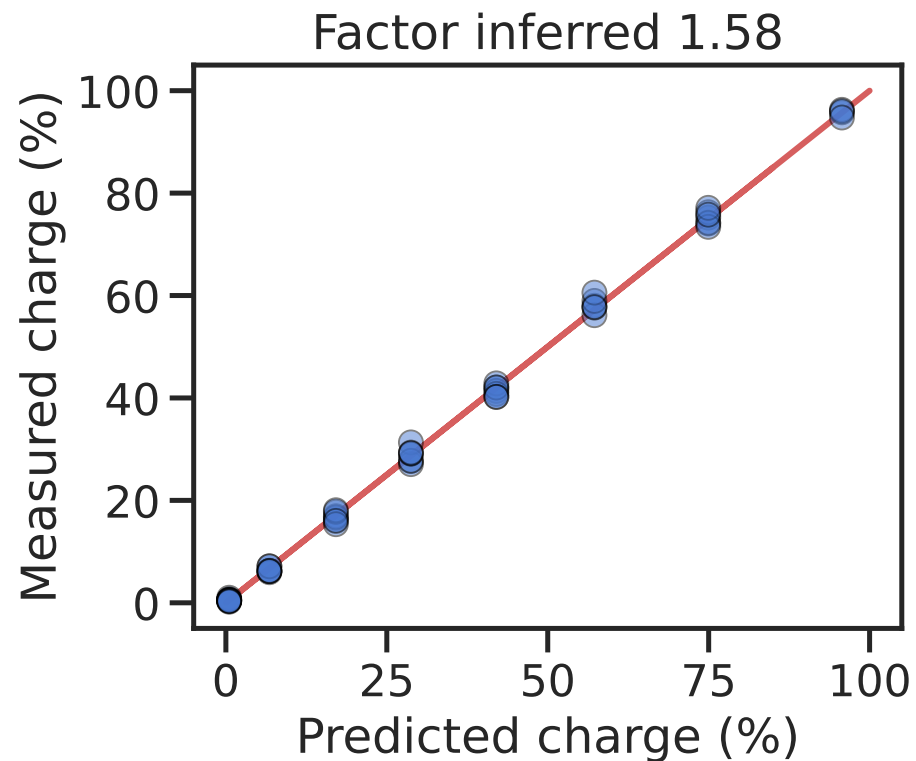
Val-CAC-2-1



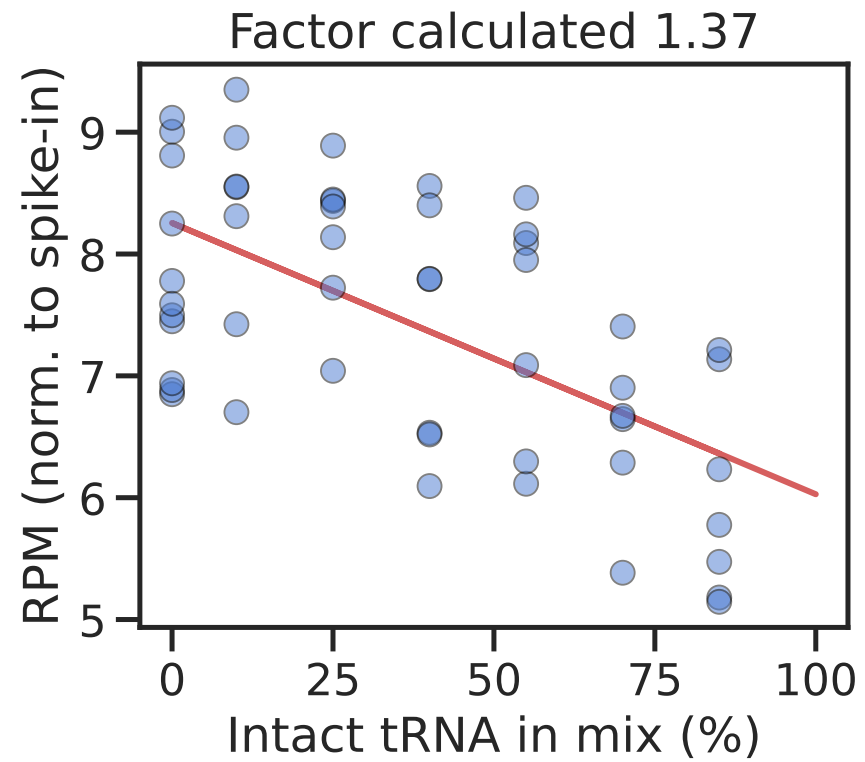
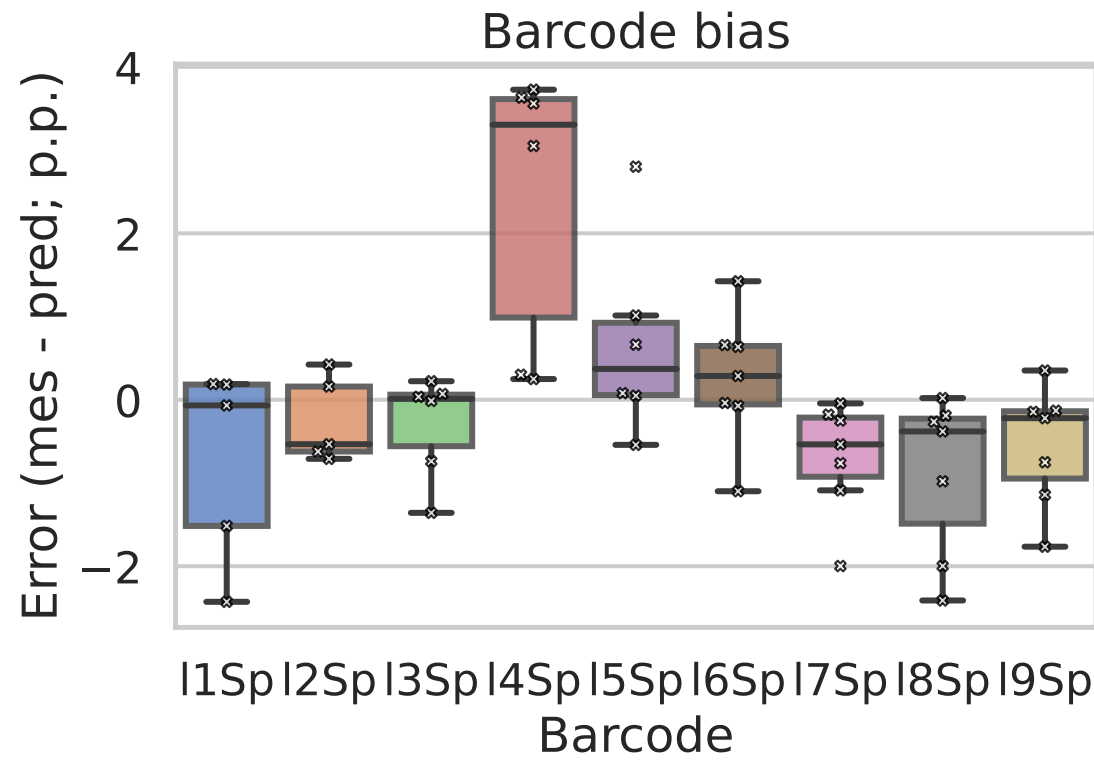
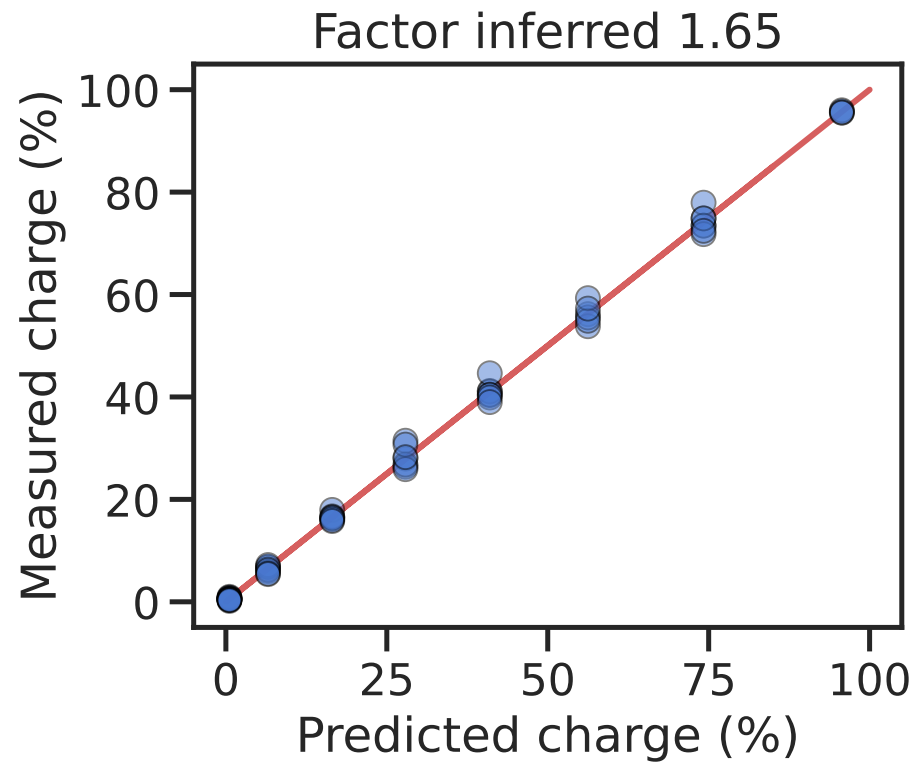
Arg-CCG-1-3



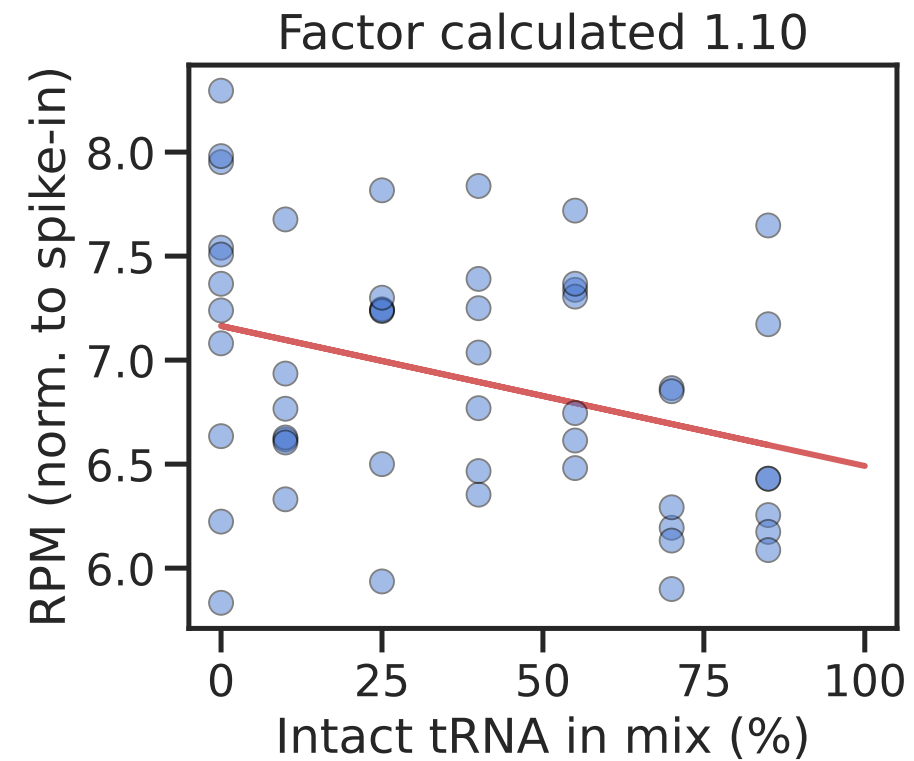
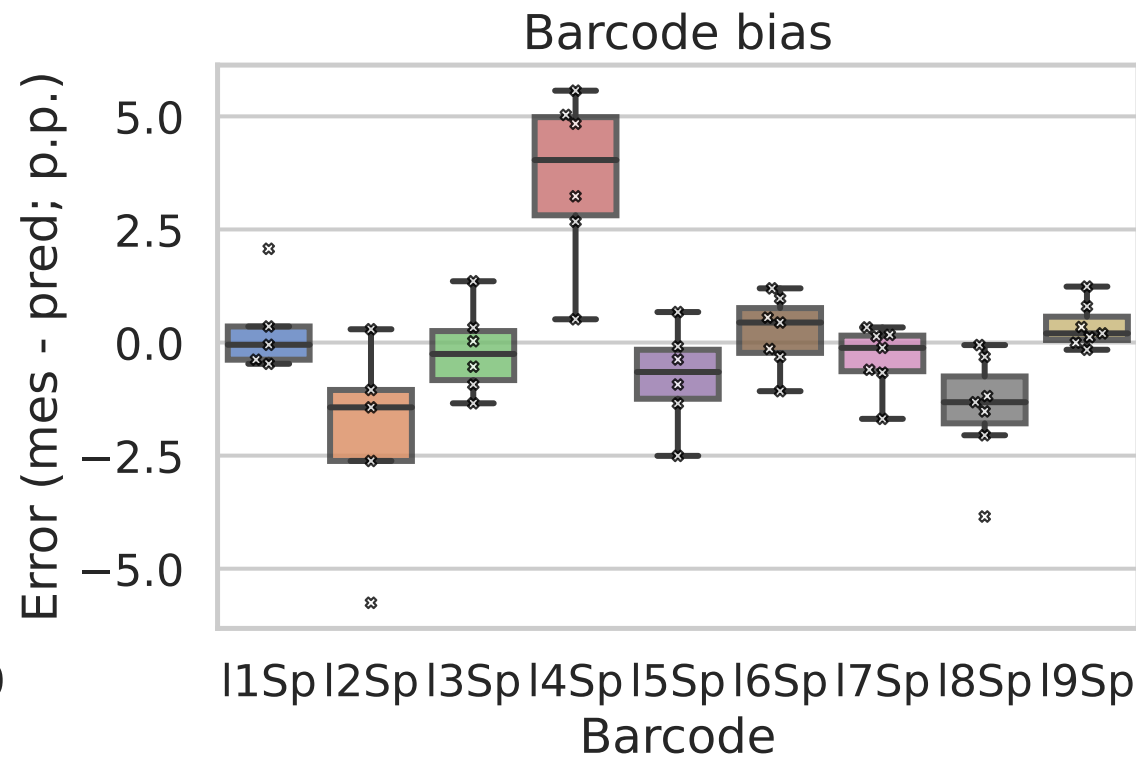
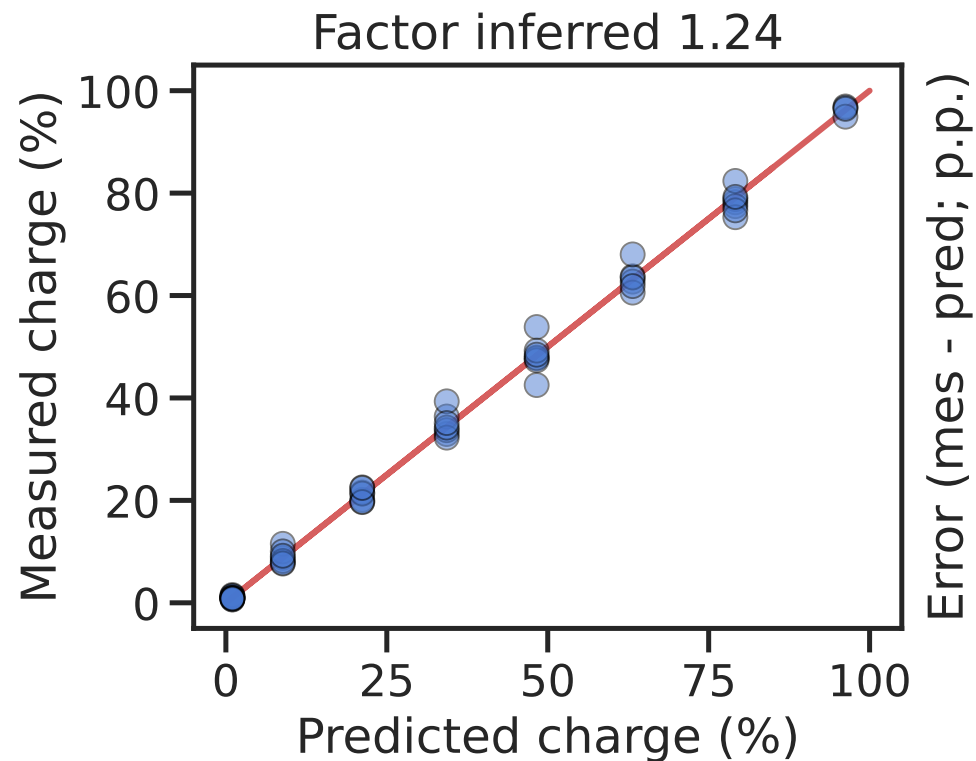
Arg-CCT-2-1



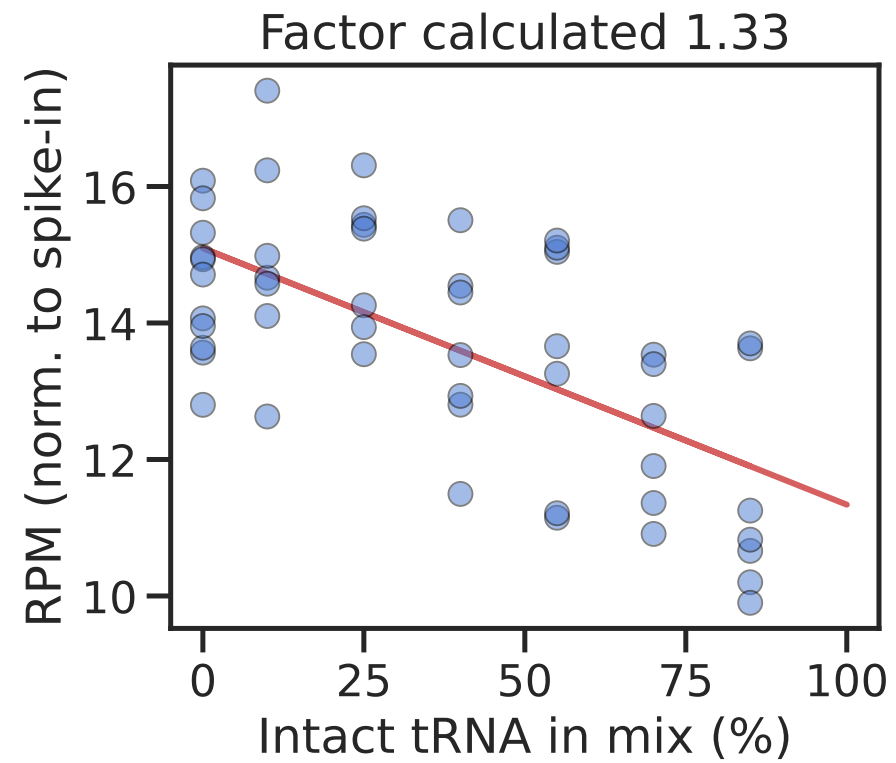
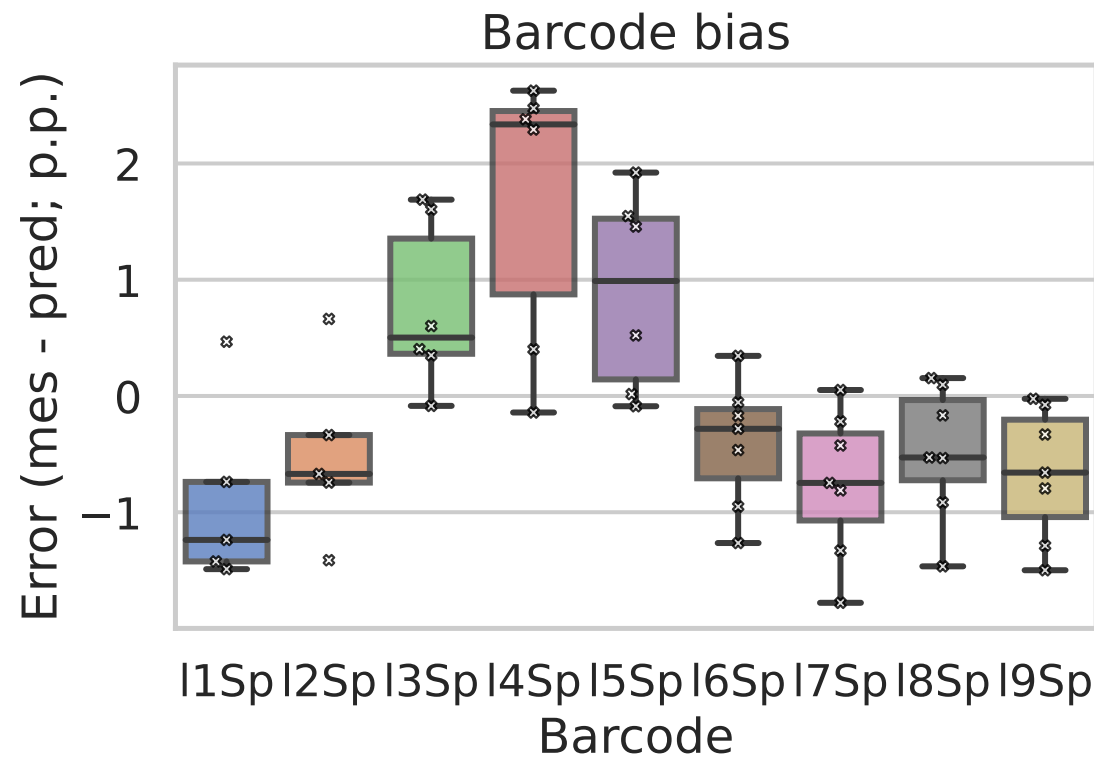
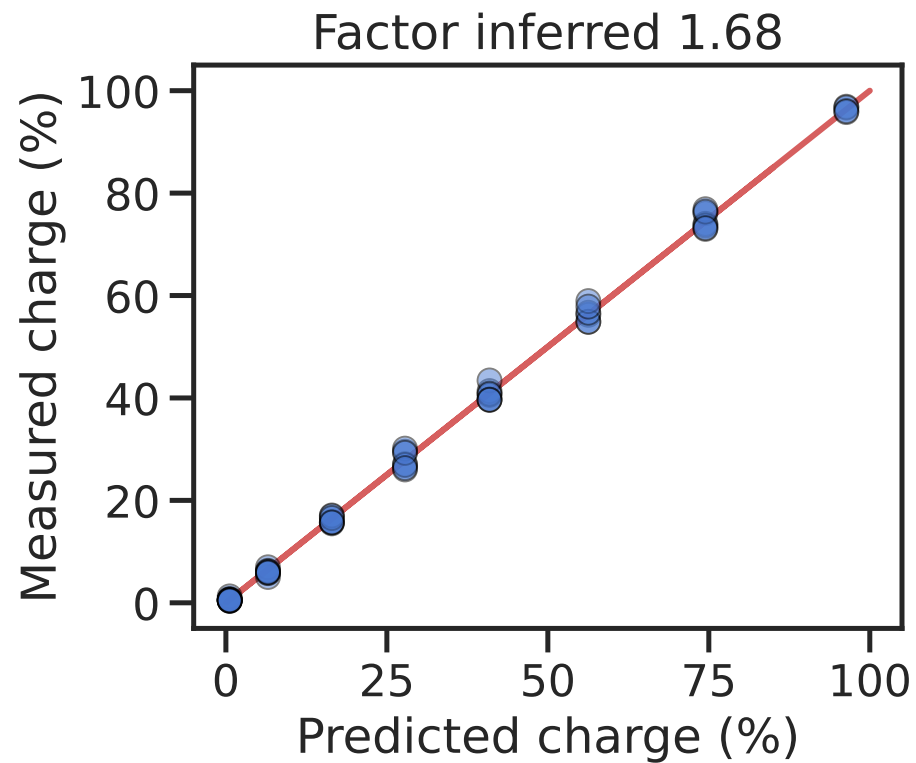
Arg-CCT-3-1



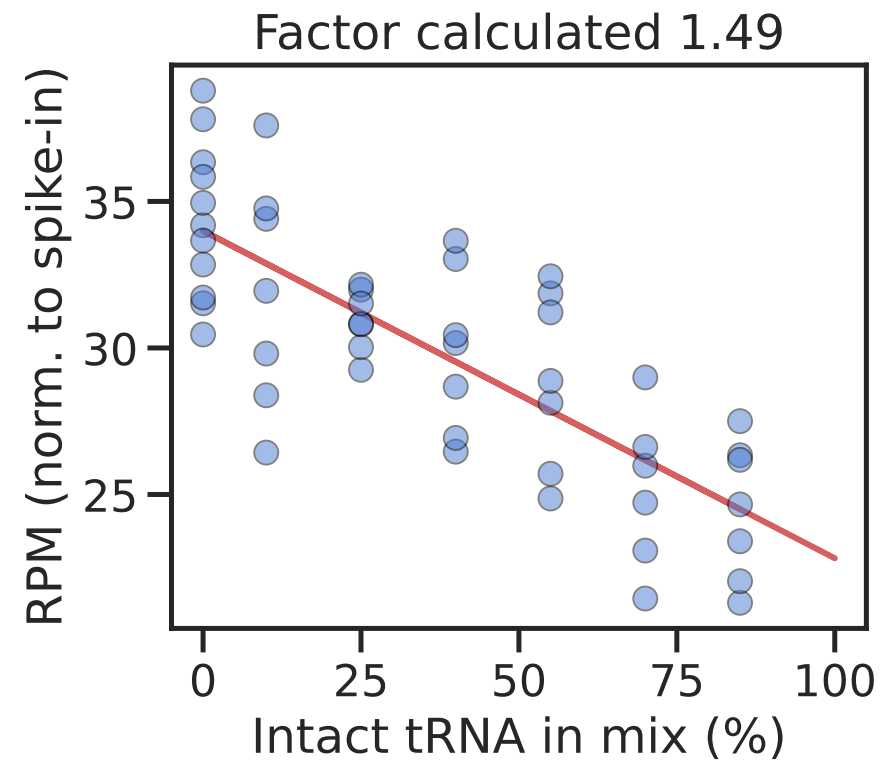
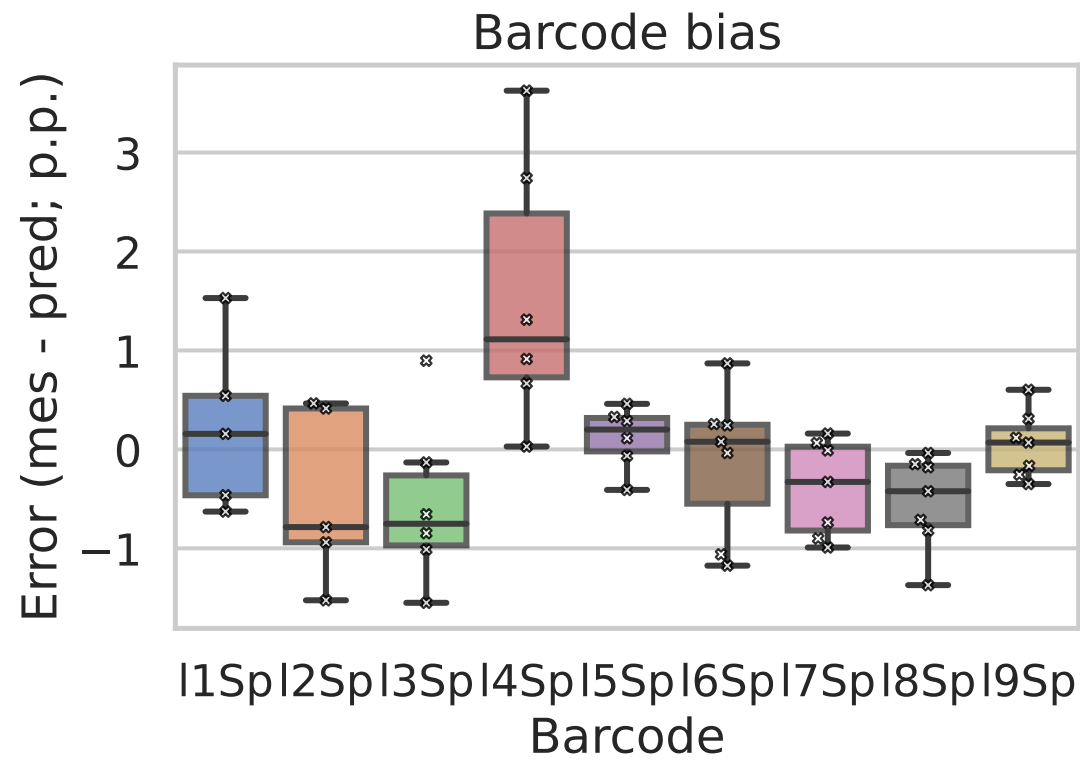
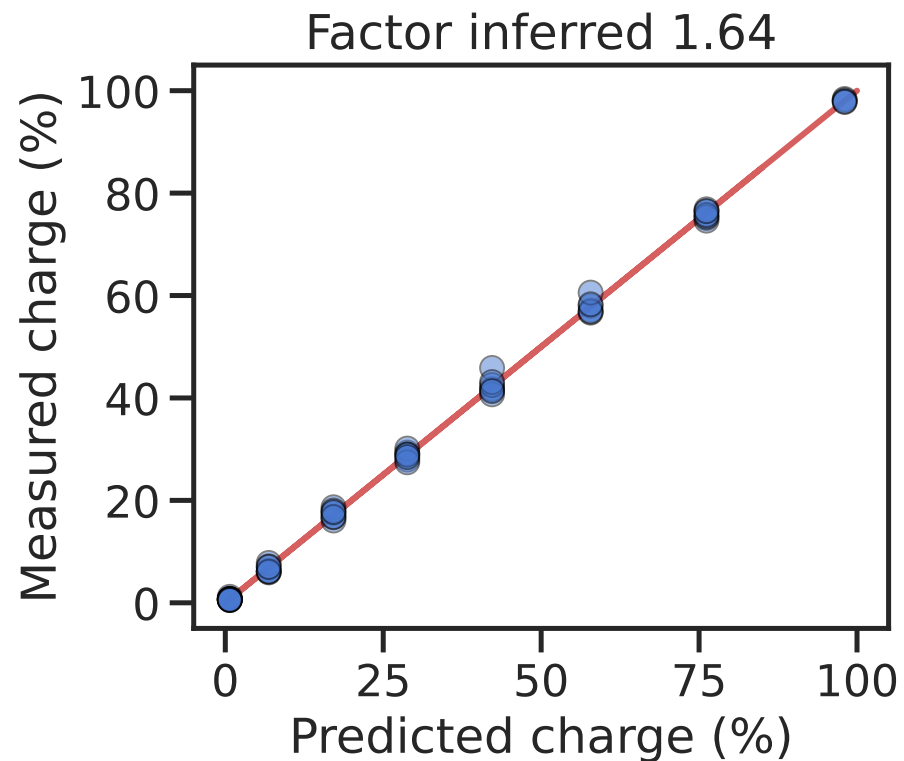
Cys-GCA-4-1



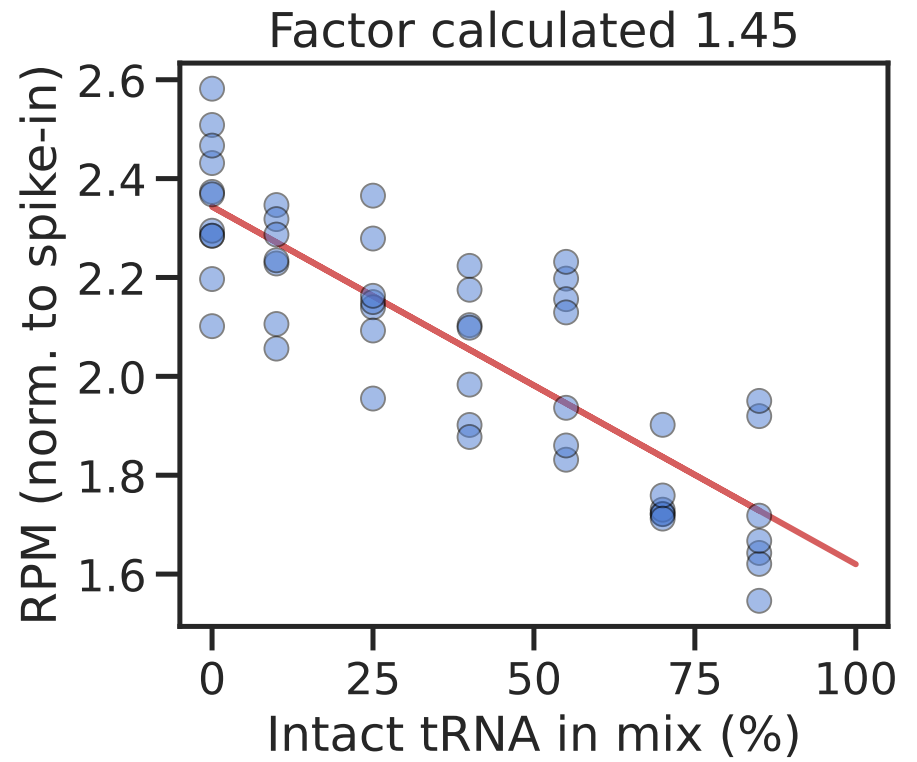
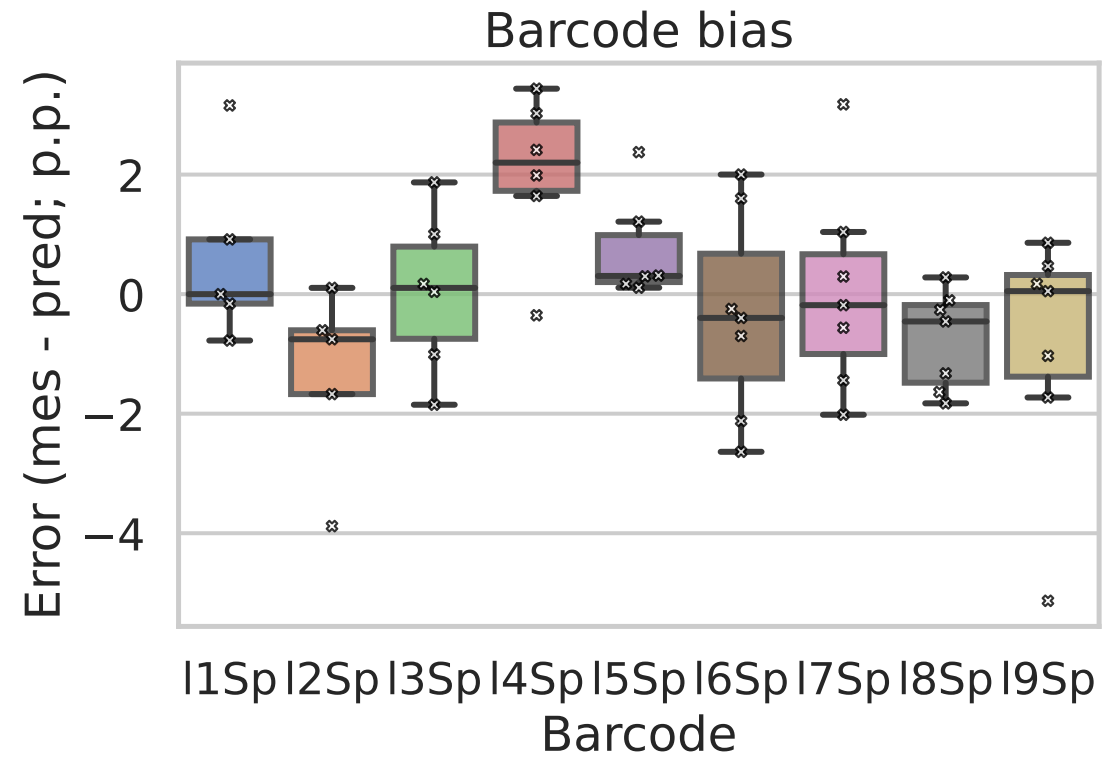
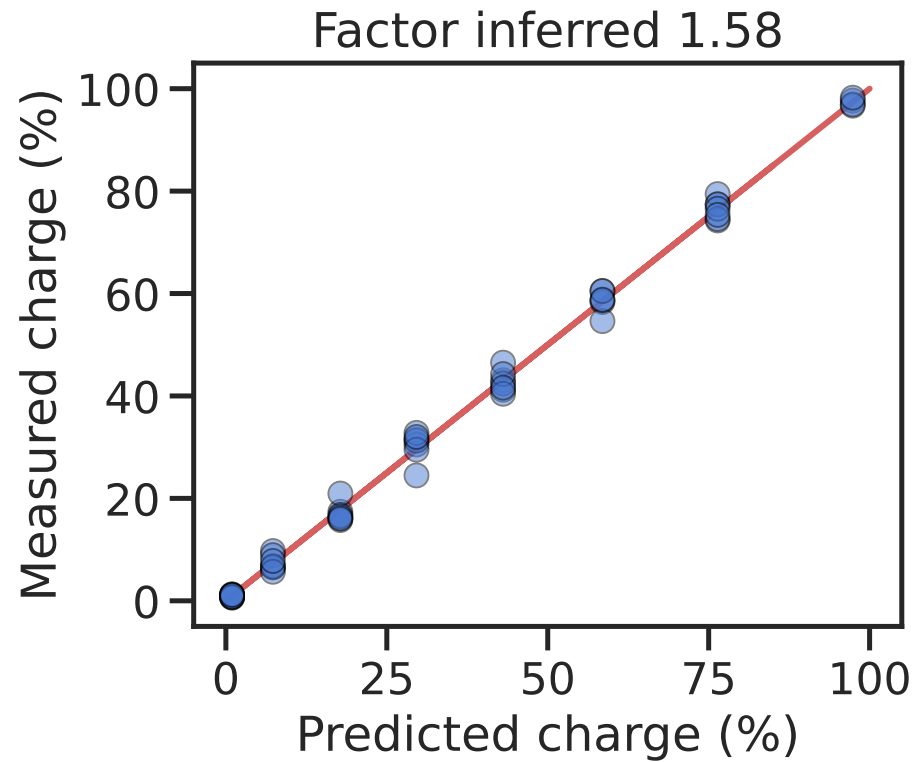
Arg-TCT-1-1



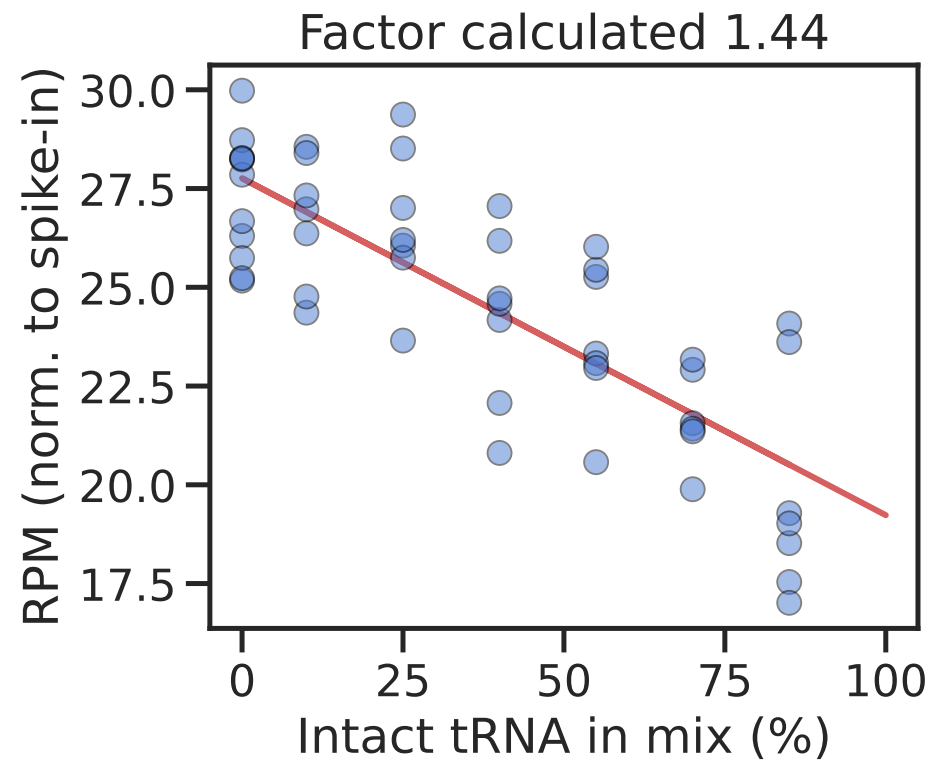
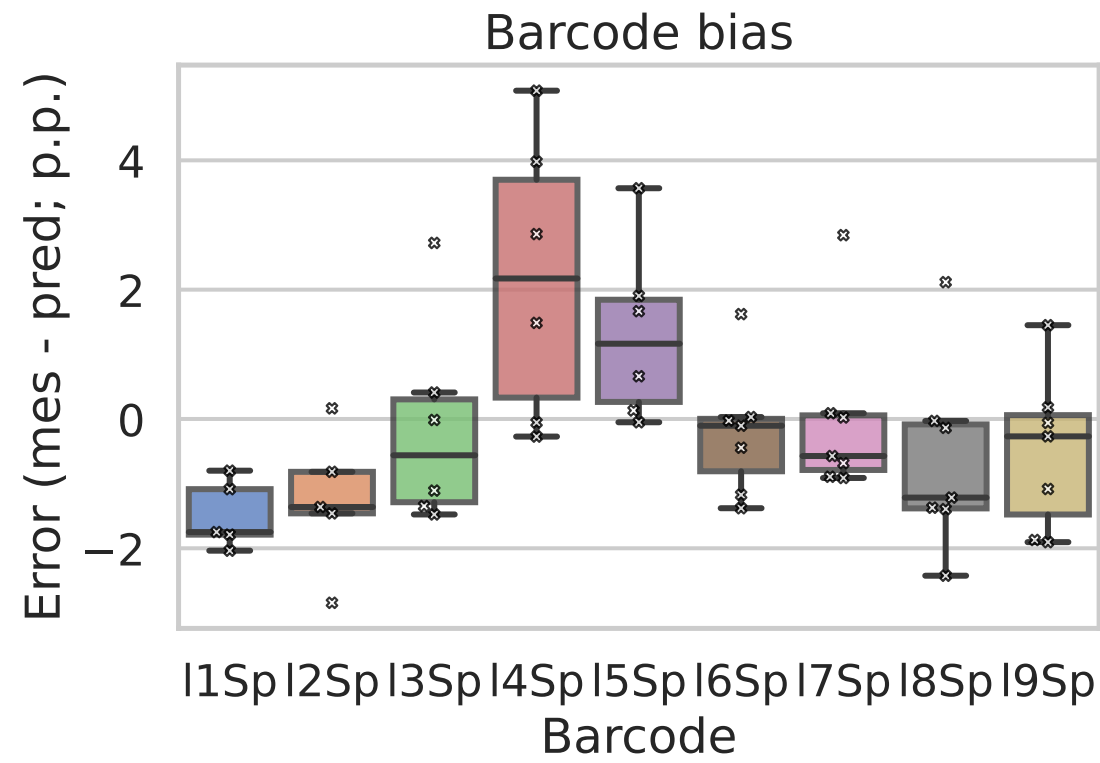
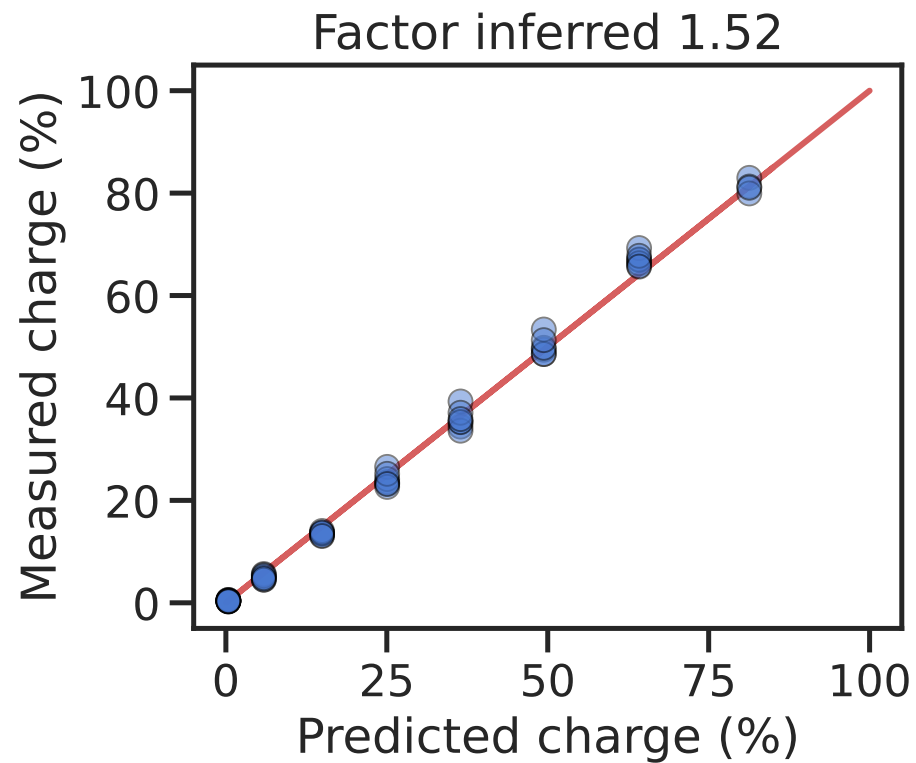
Gly-CCC-2-2



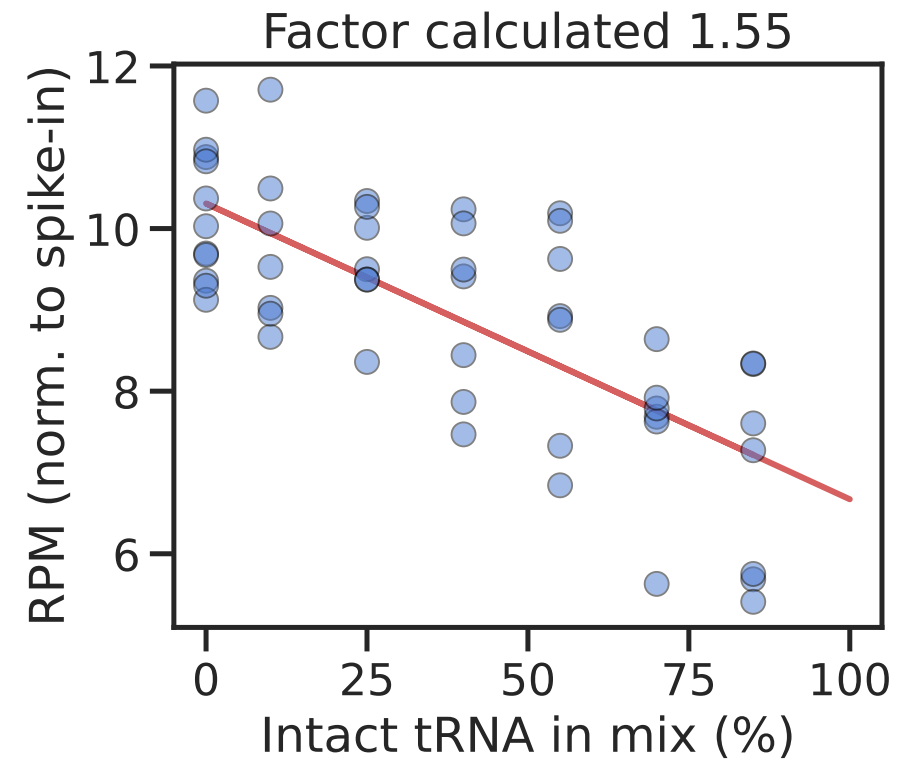
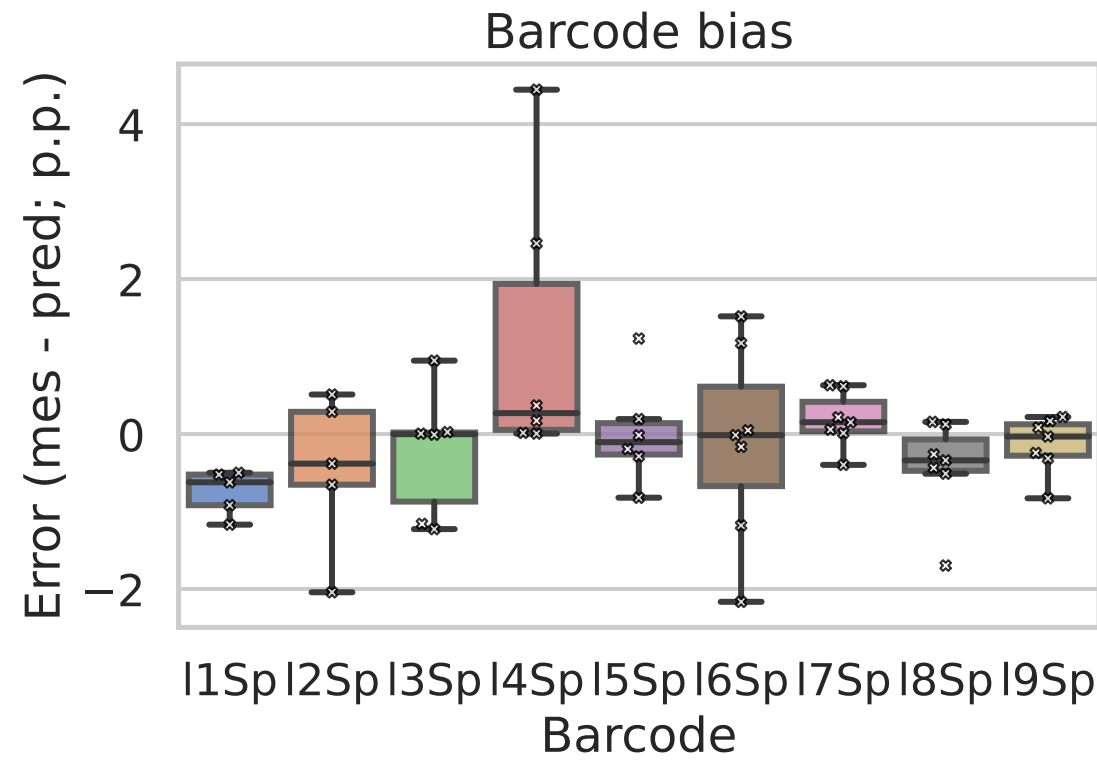
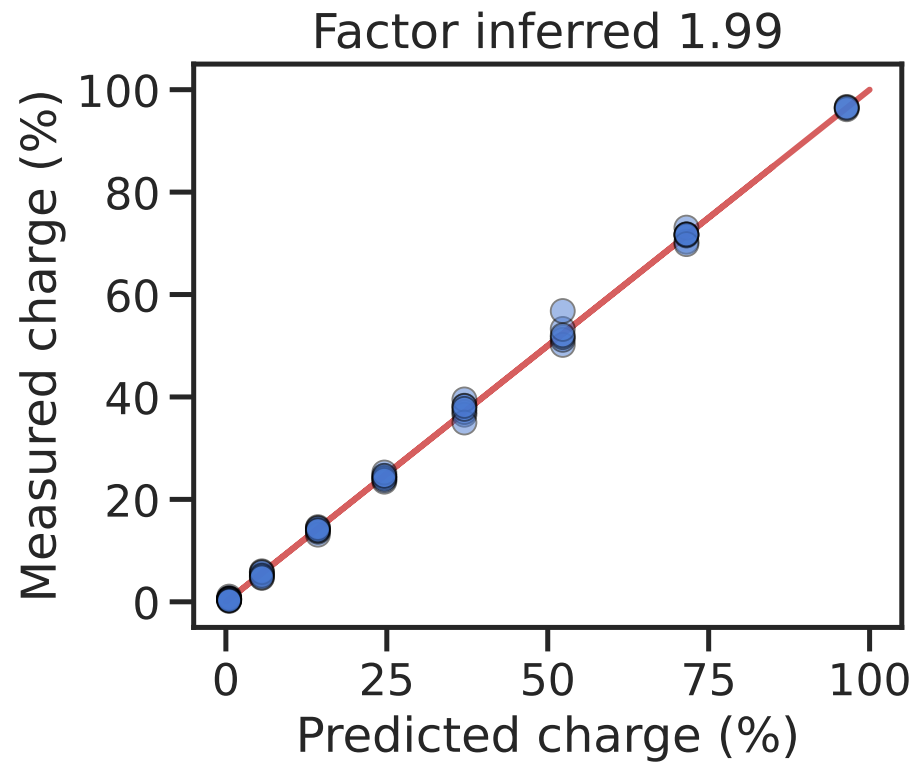
Val-CAC-4-1



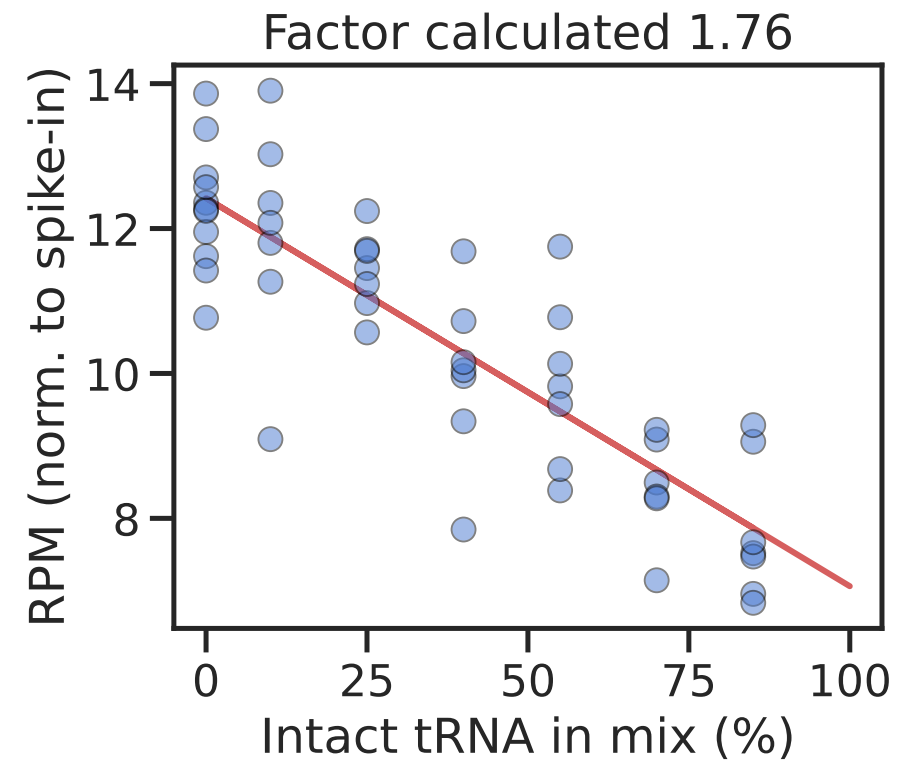
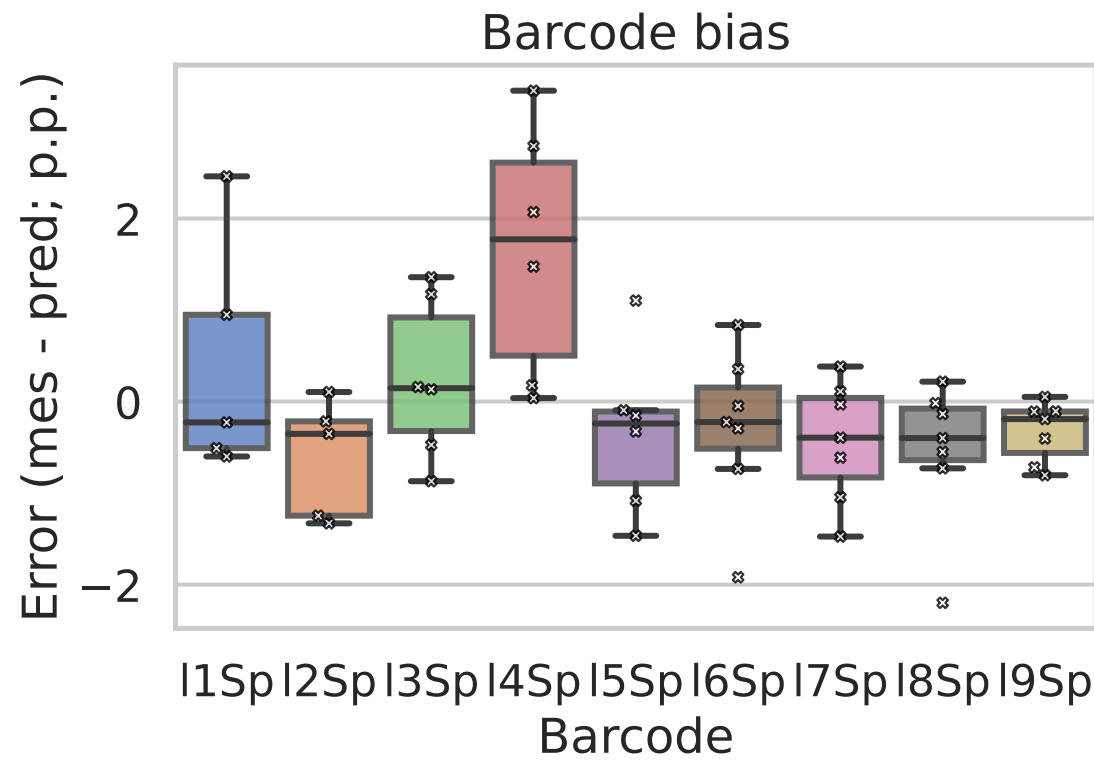
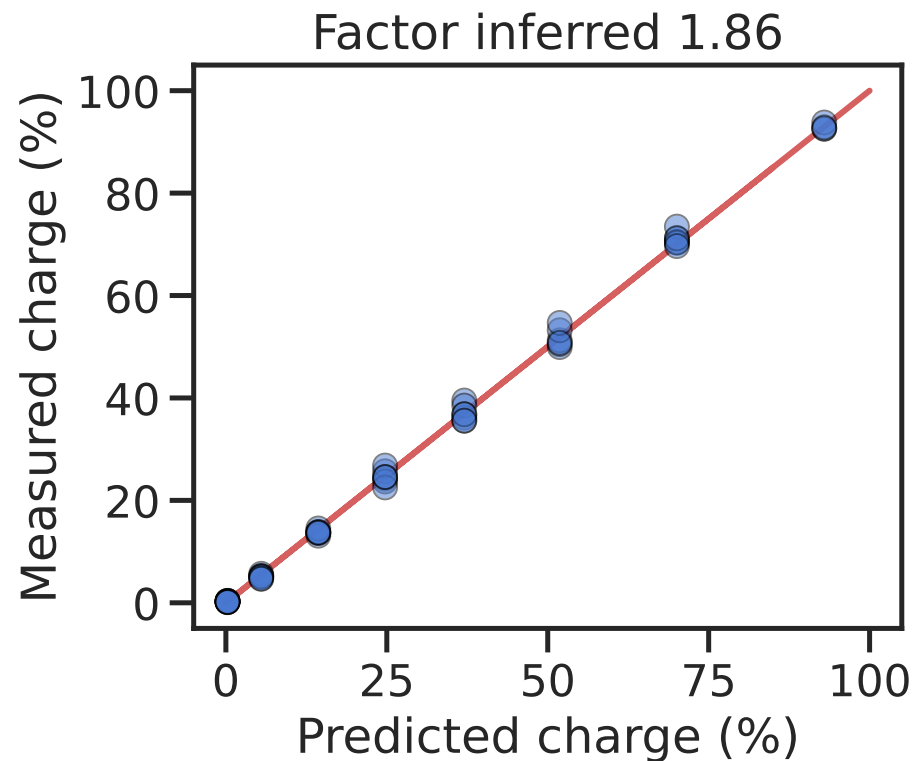
Ser-GCT-3-1



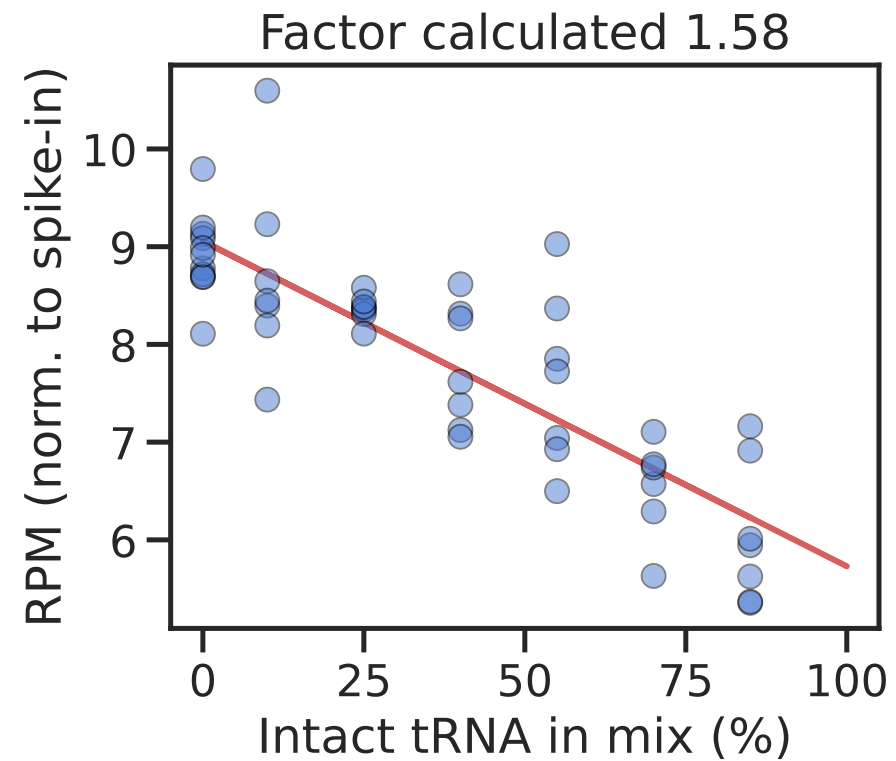
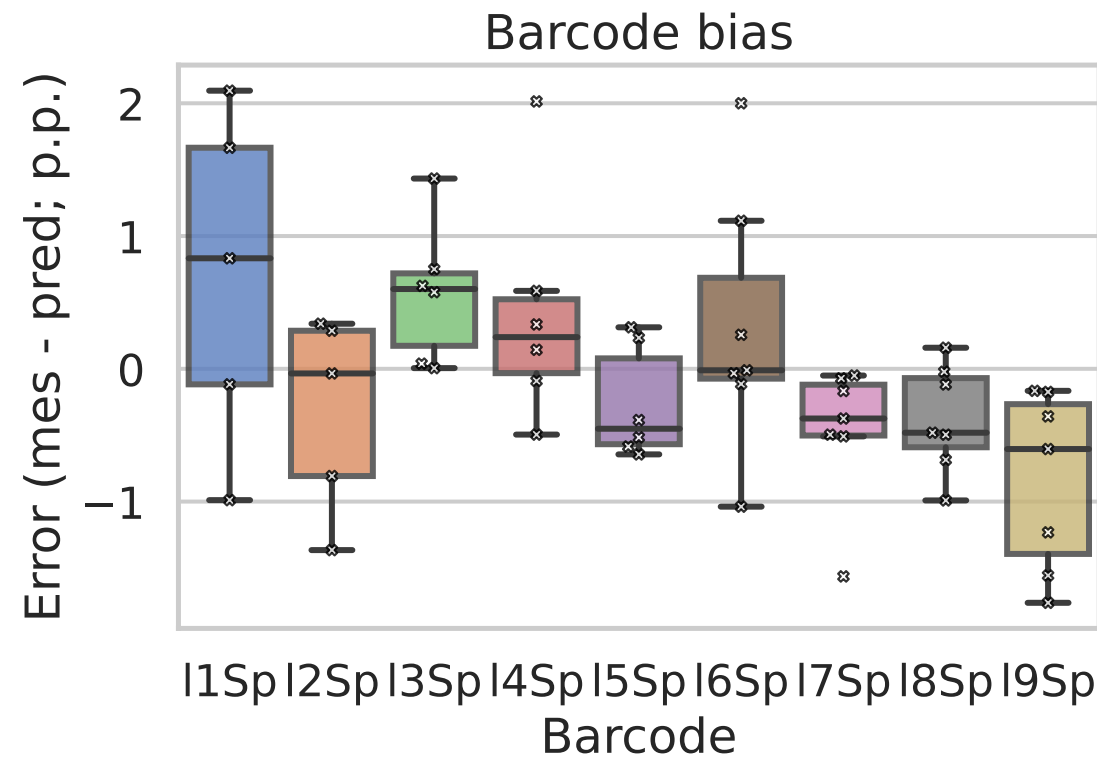
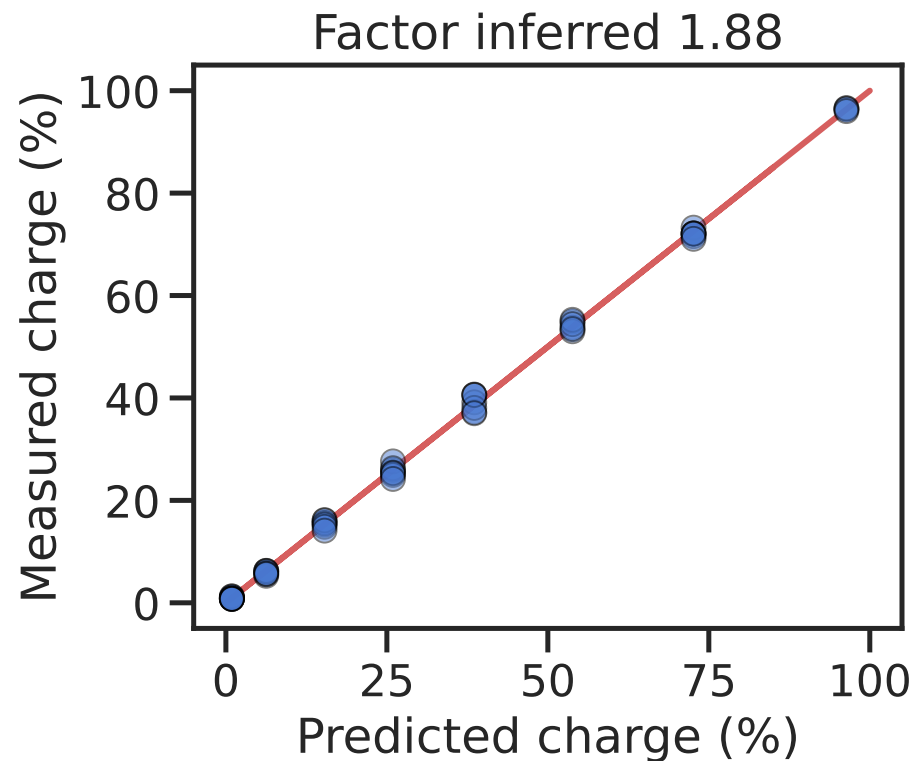
Leu-TAG-2-1



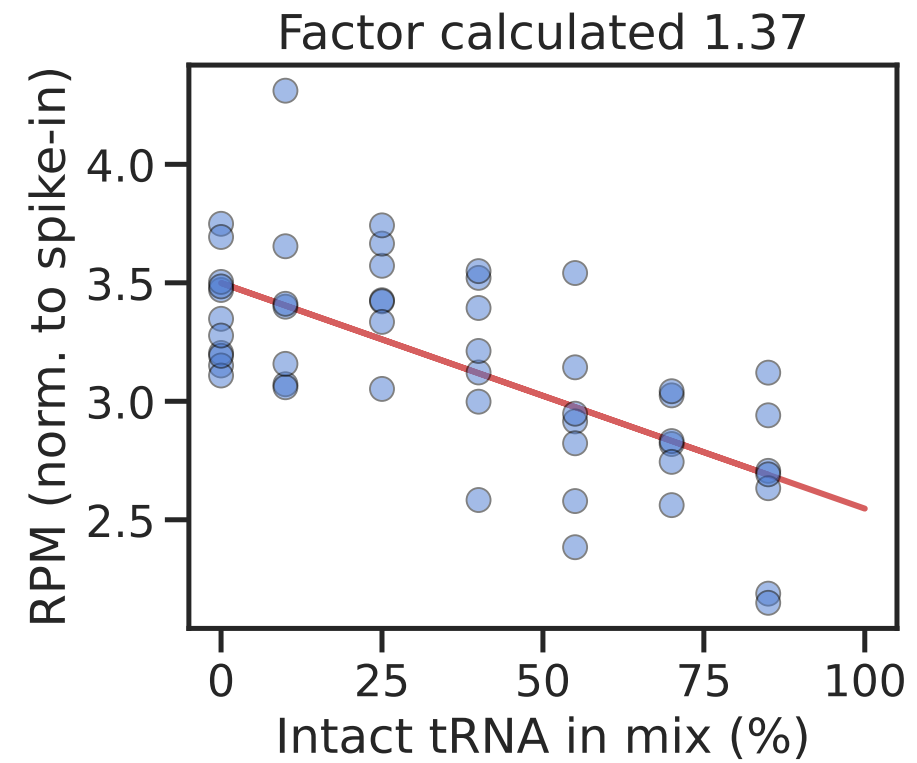
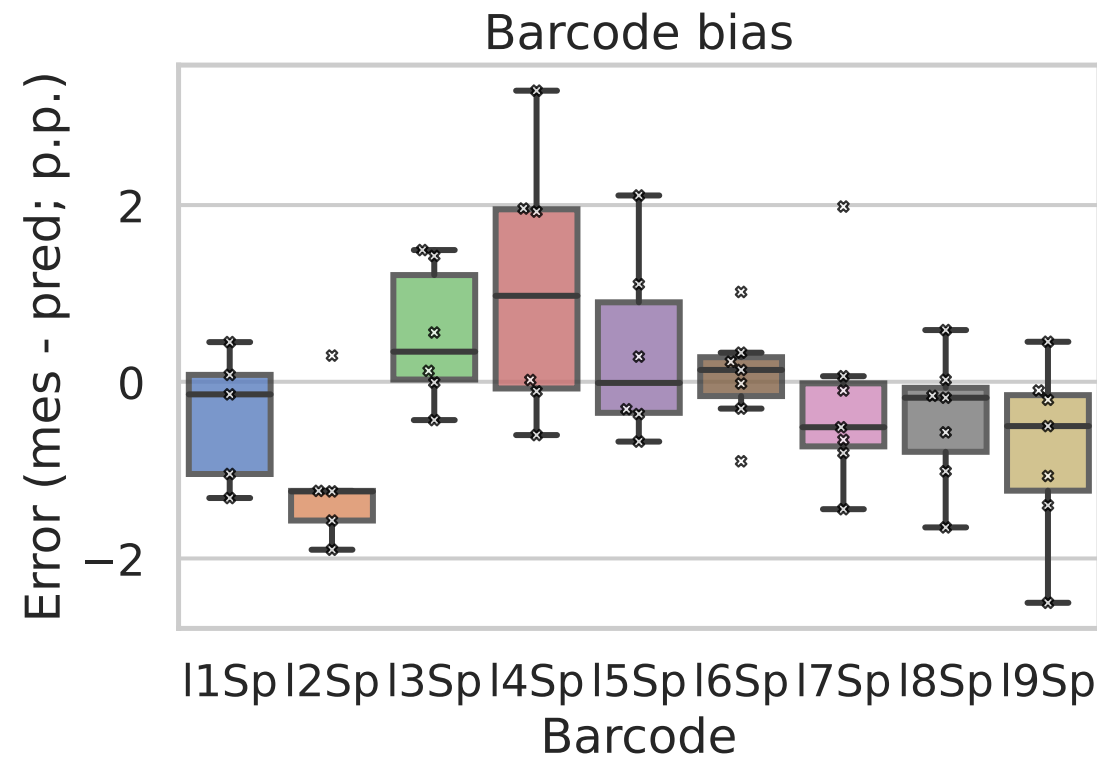
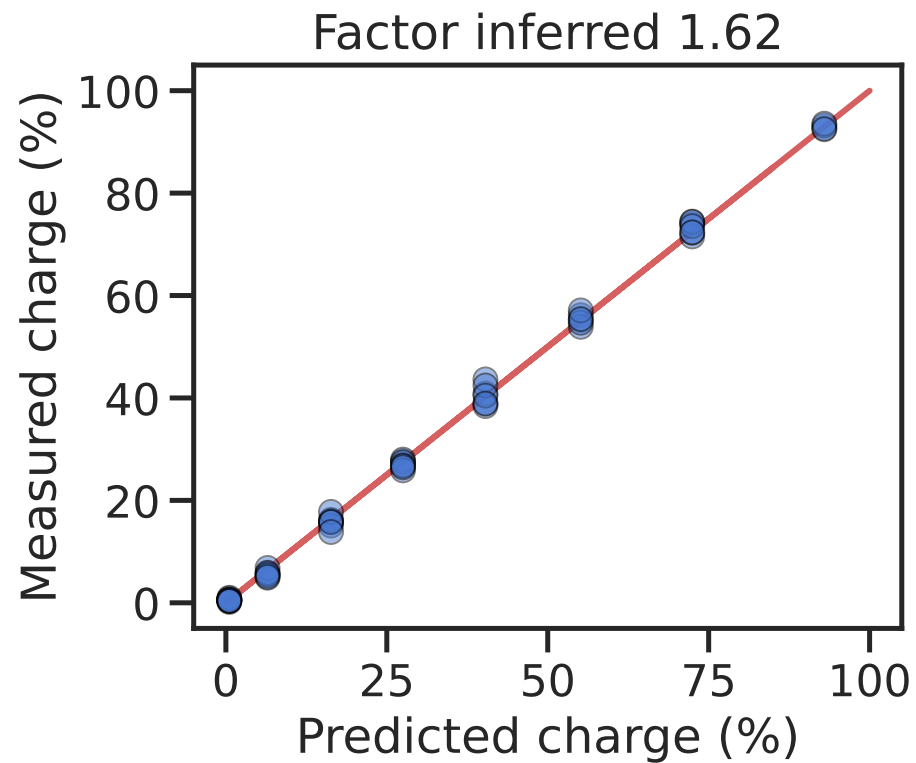
Pro-TGG-2-1



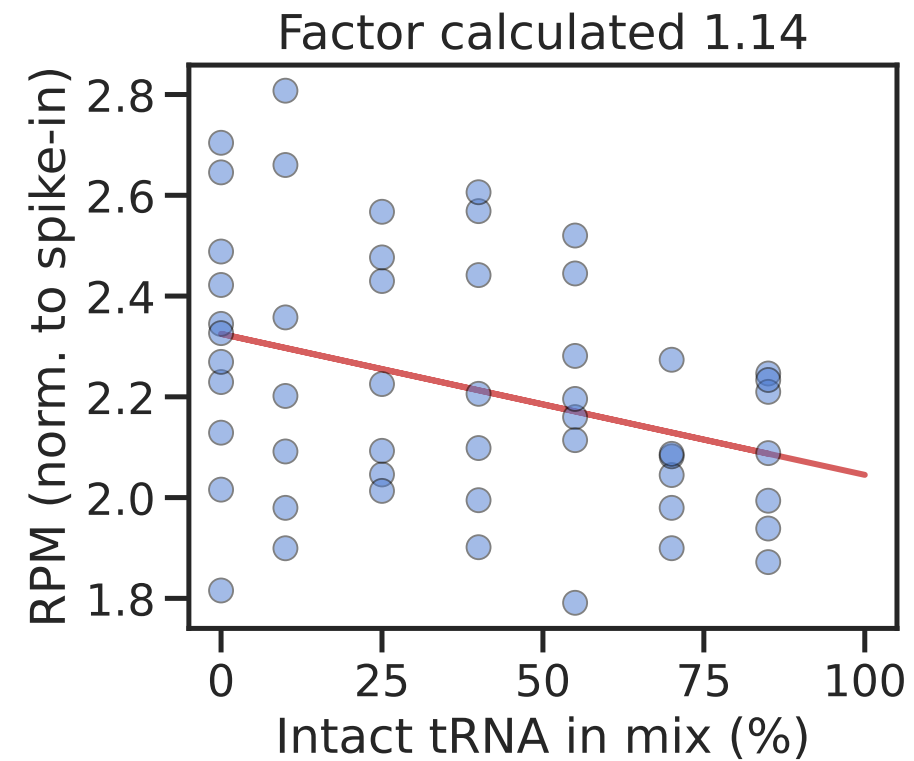
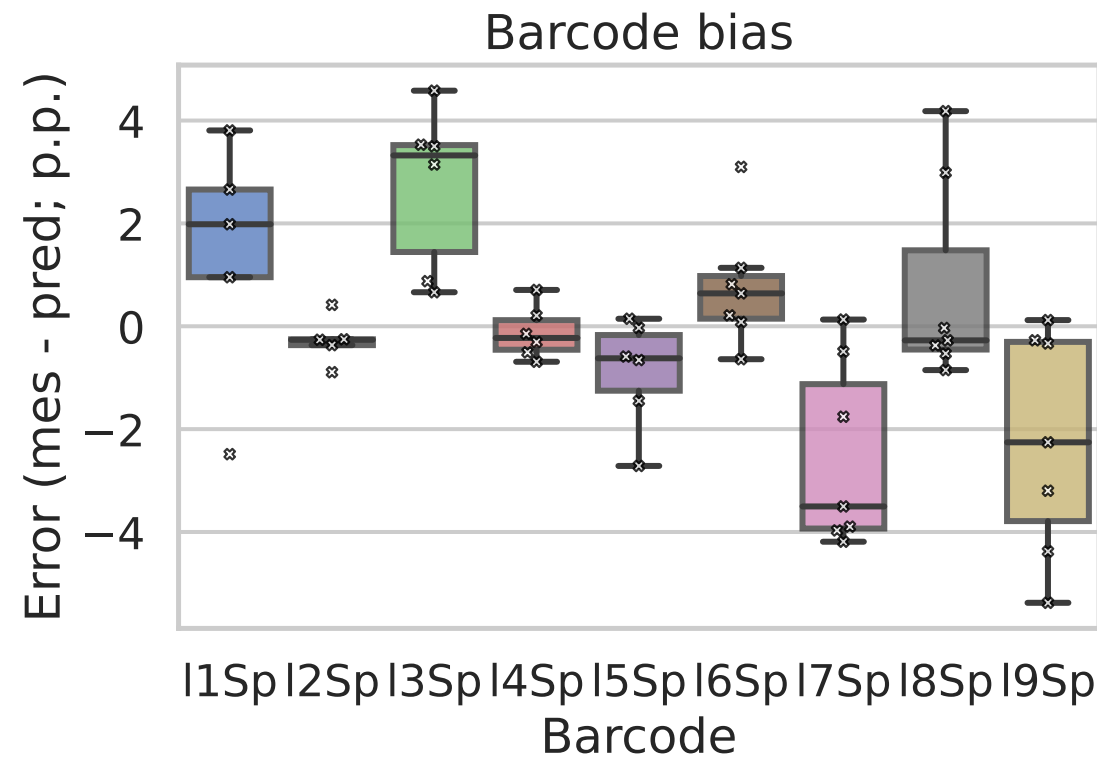
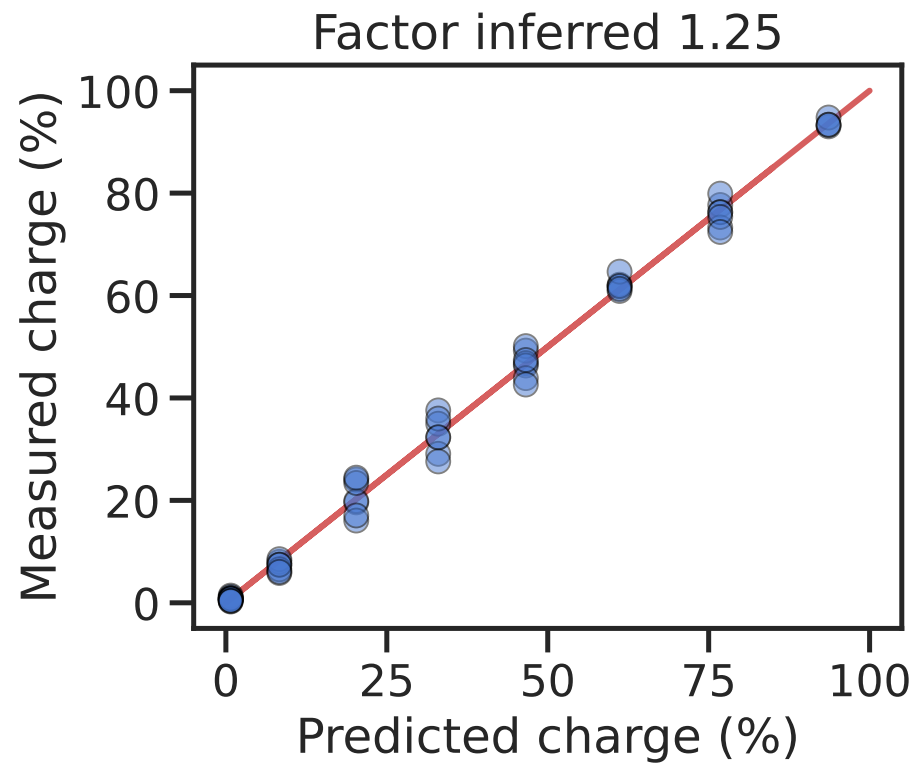
Ile-AAT-4-1



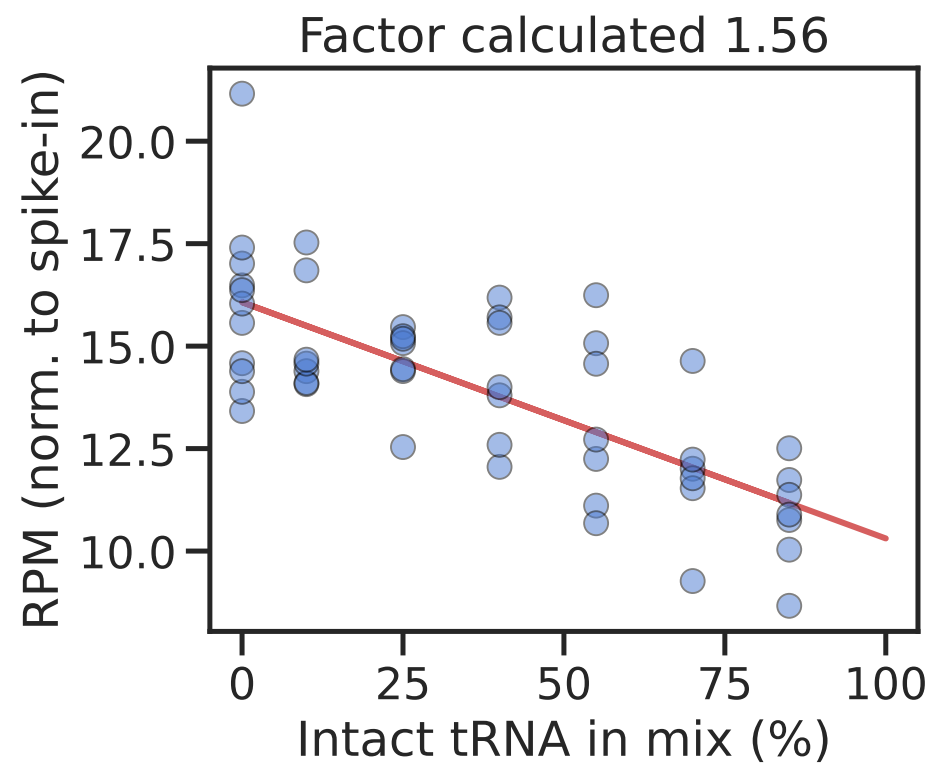
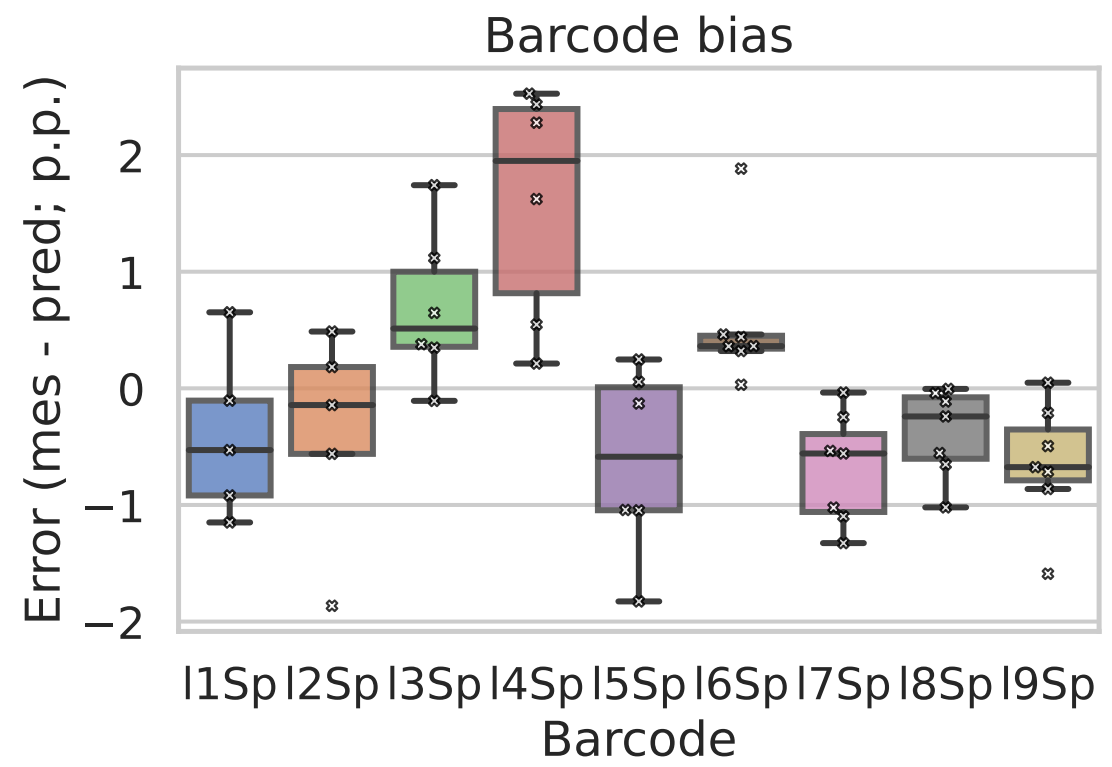
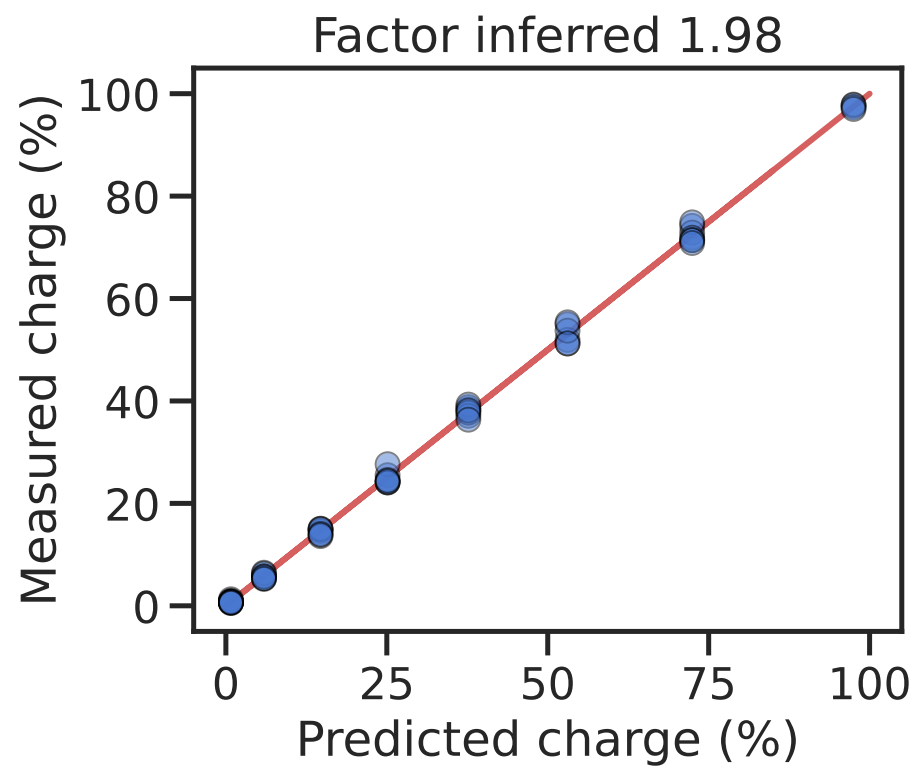
mito-Gln-TTG



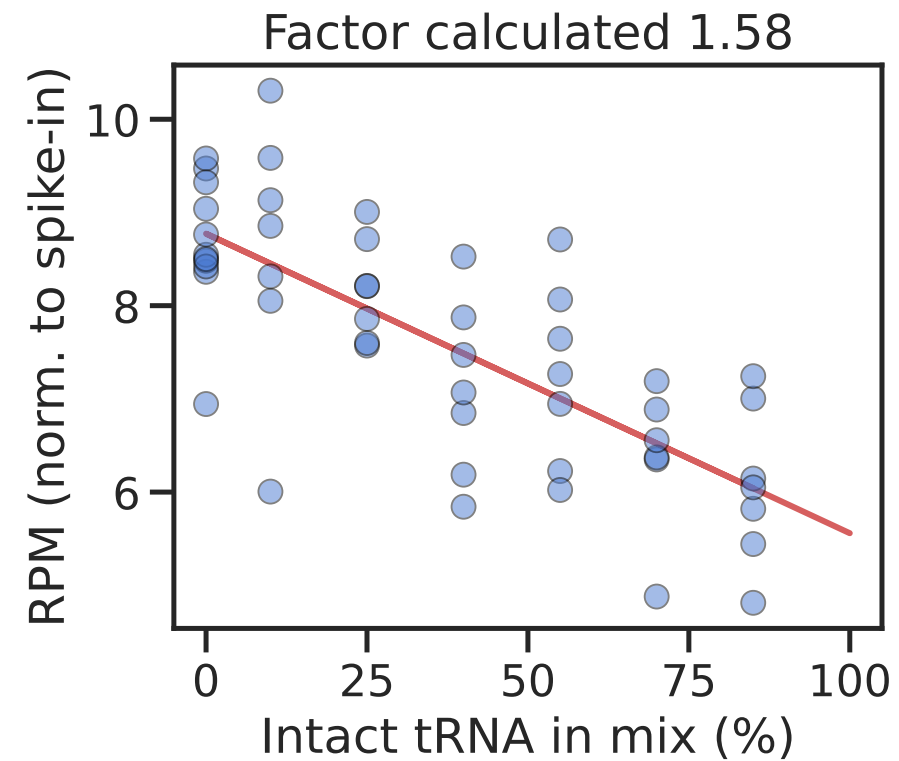
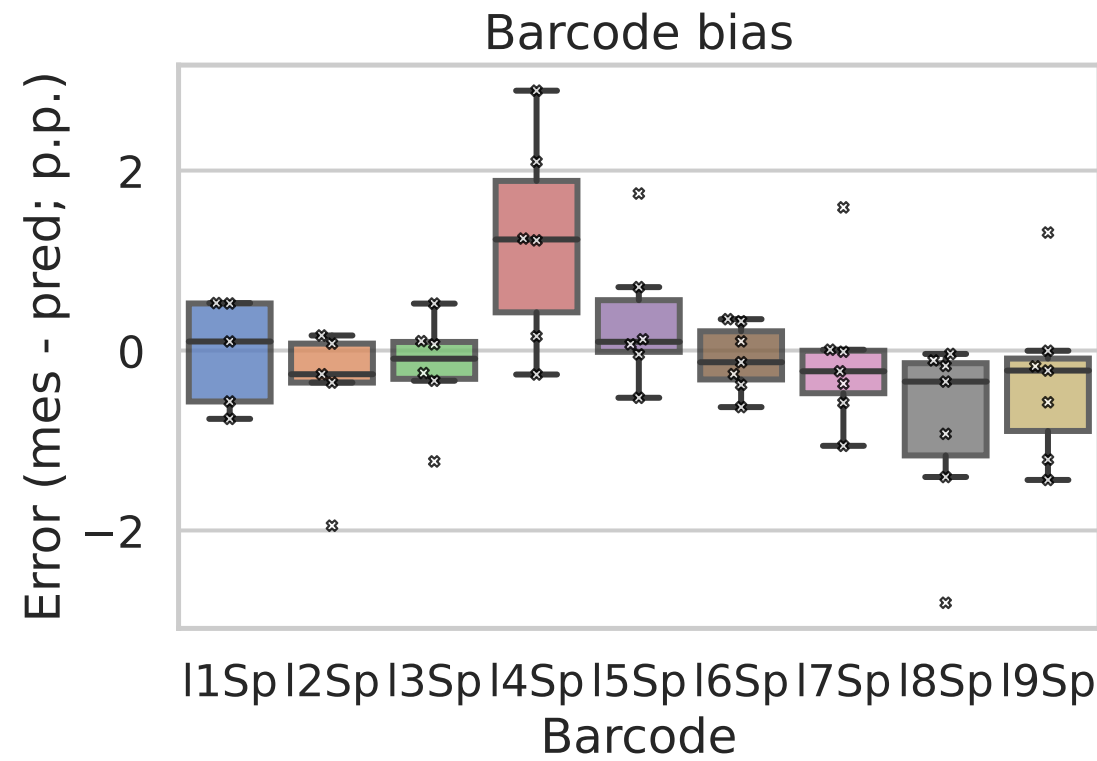
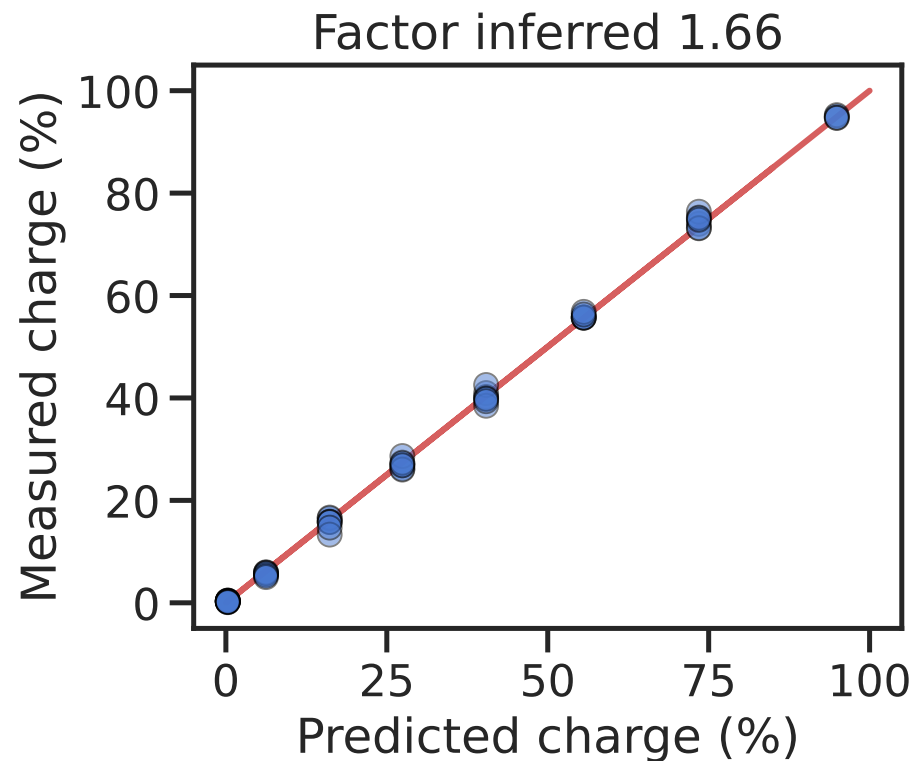
mito-Cys-GCA



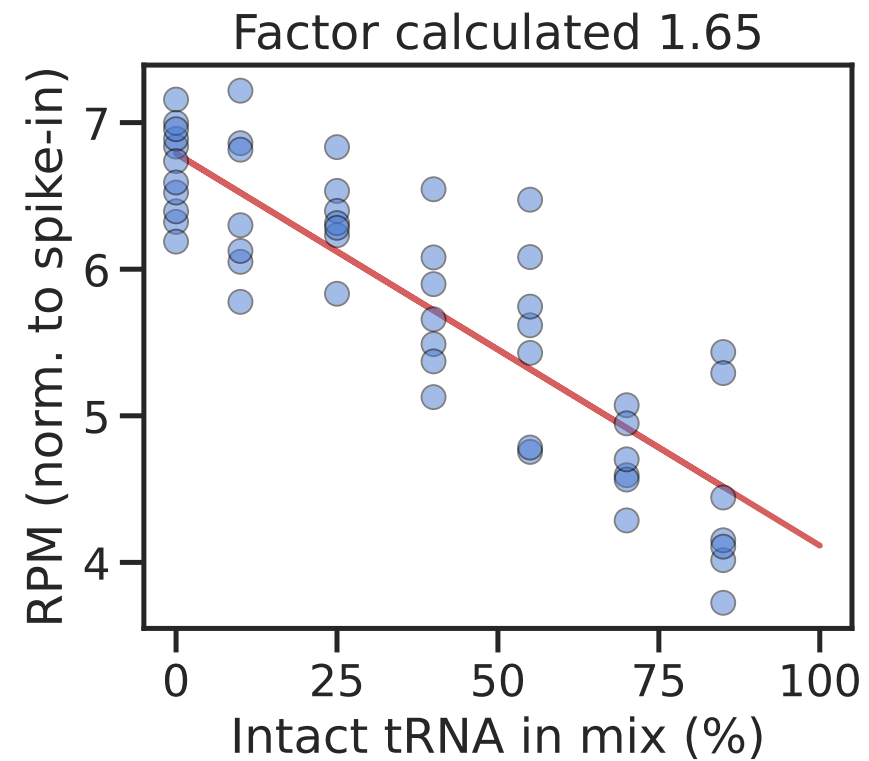
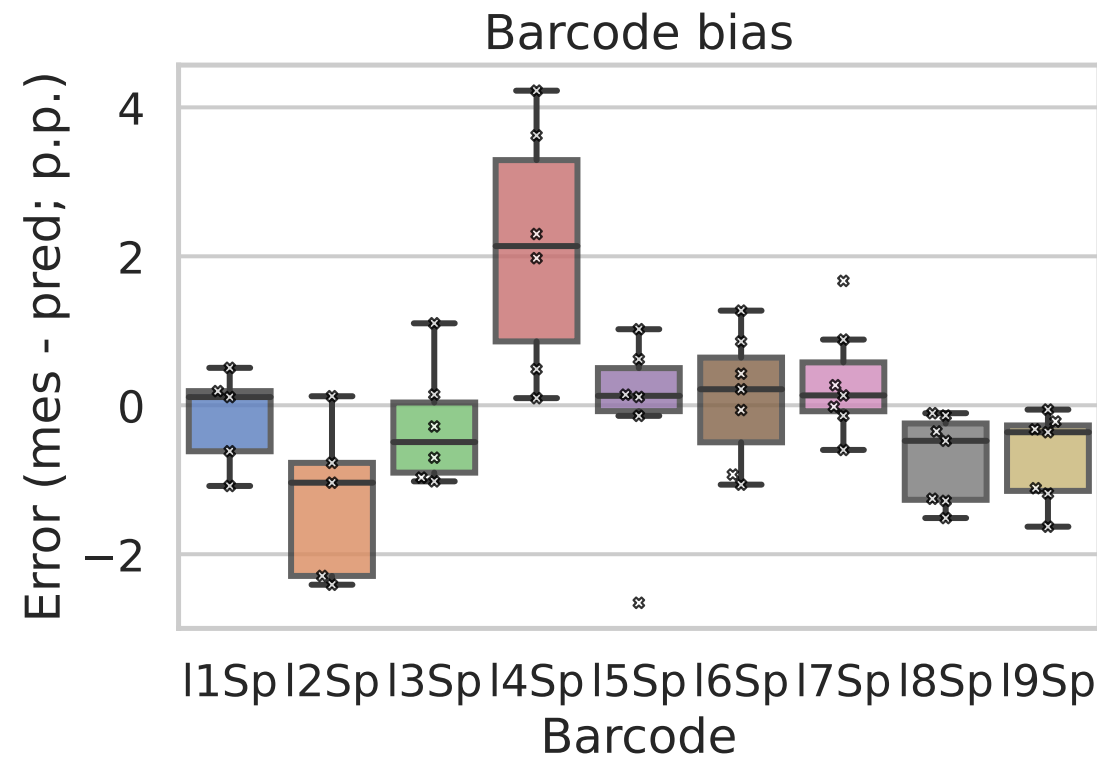
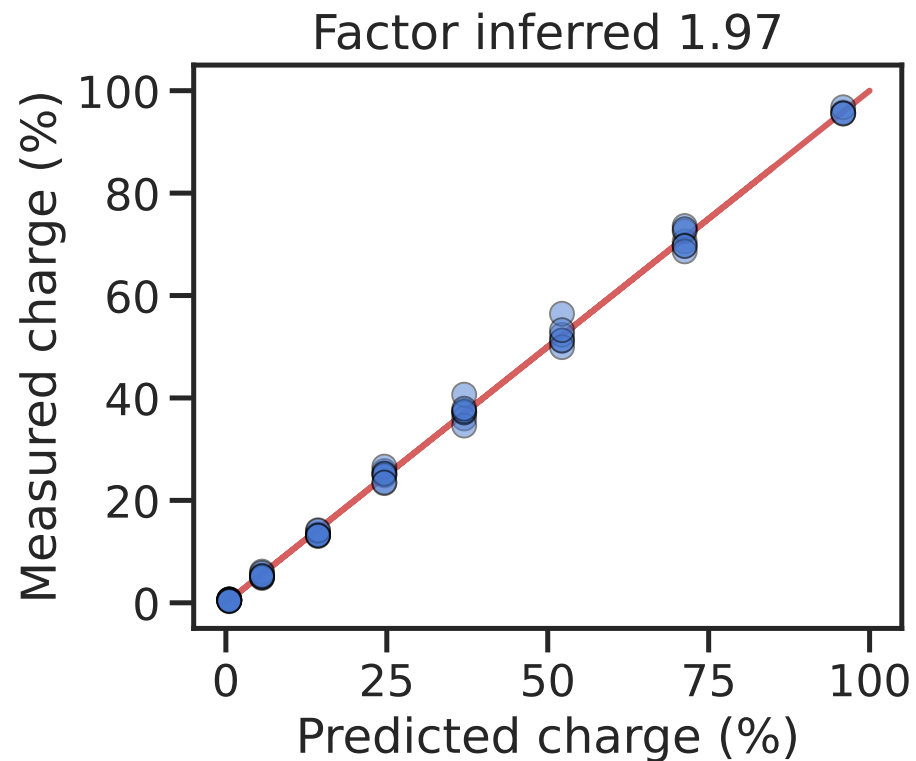
Trp-CCA-2-1



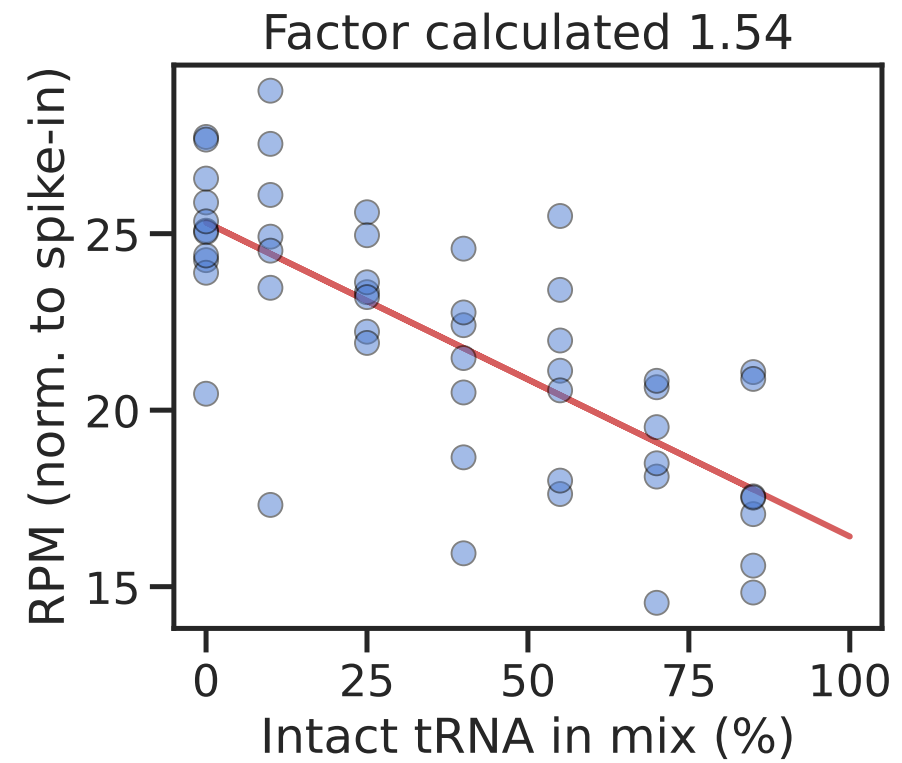
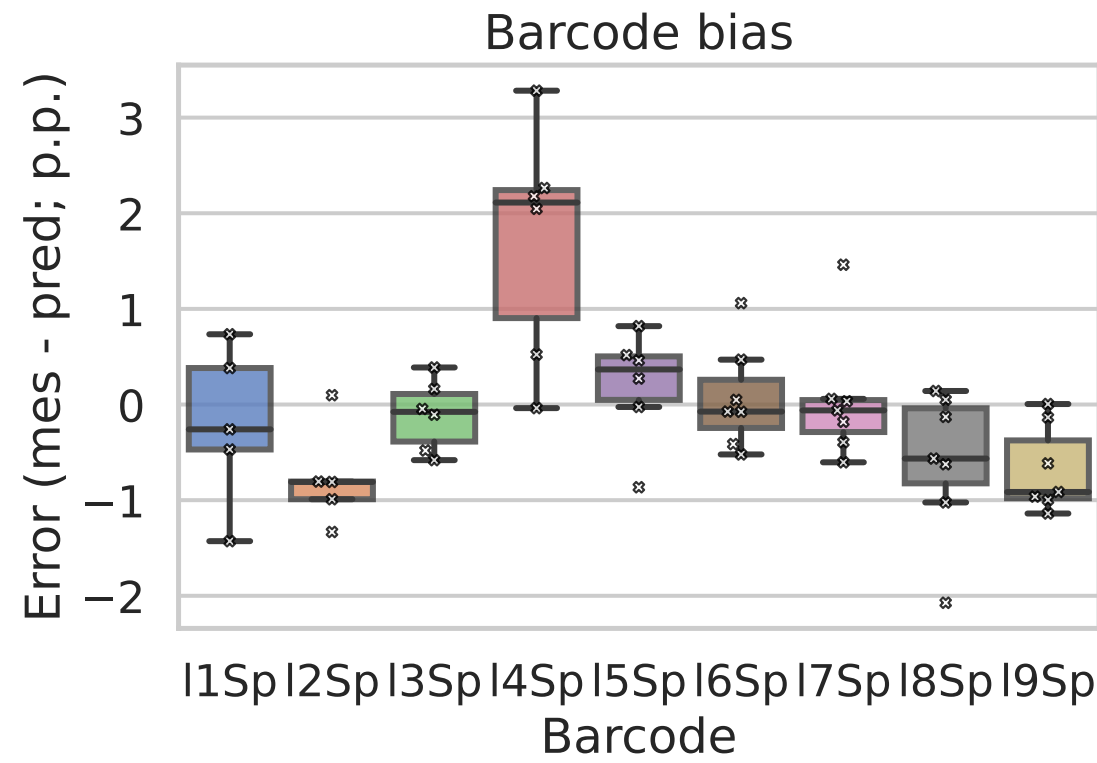
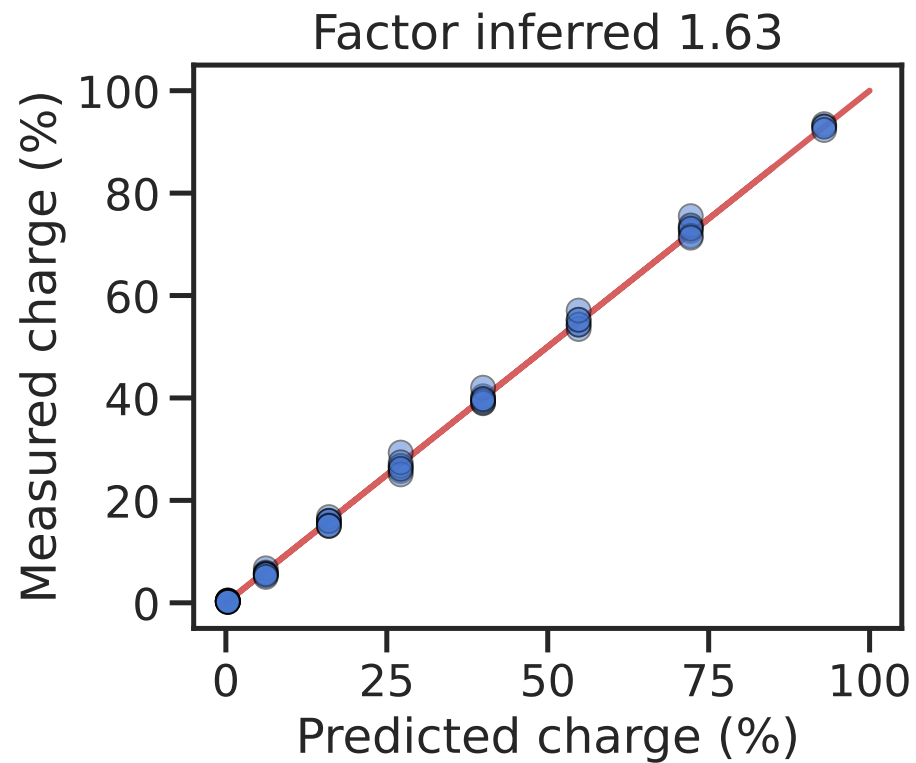
Pro-CGG-1-1



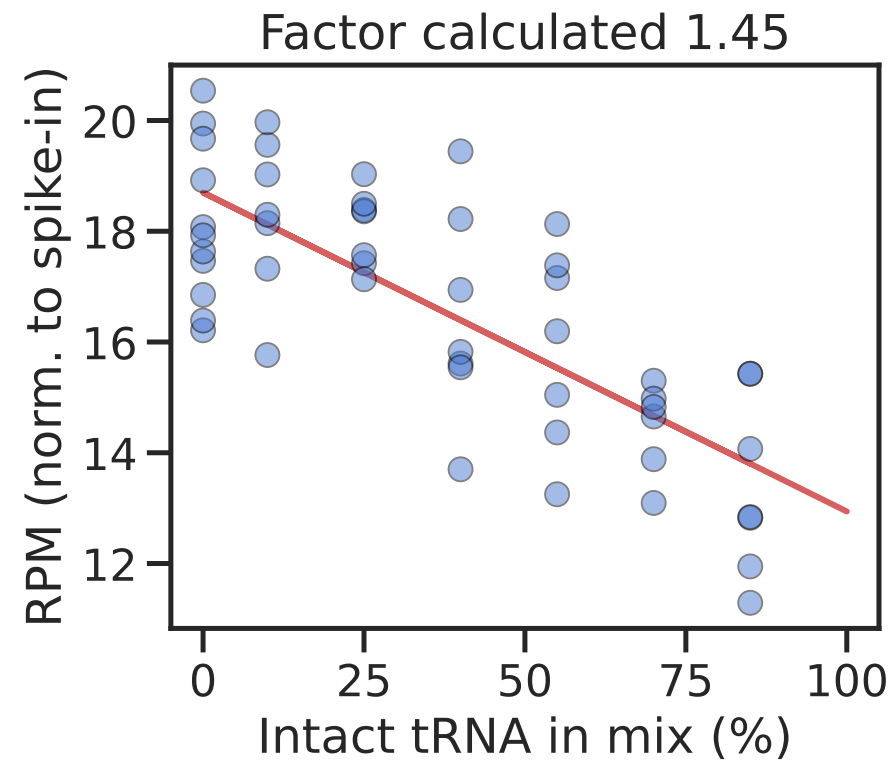
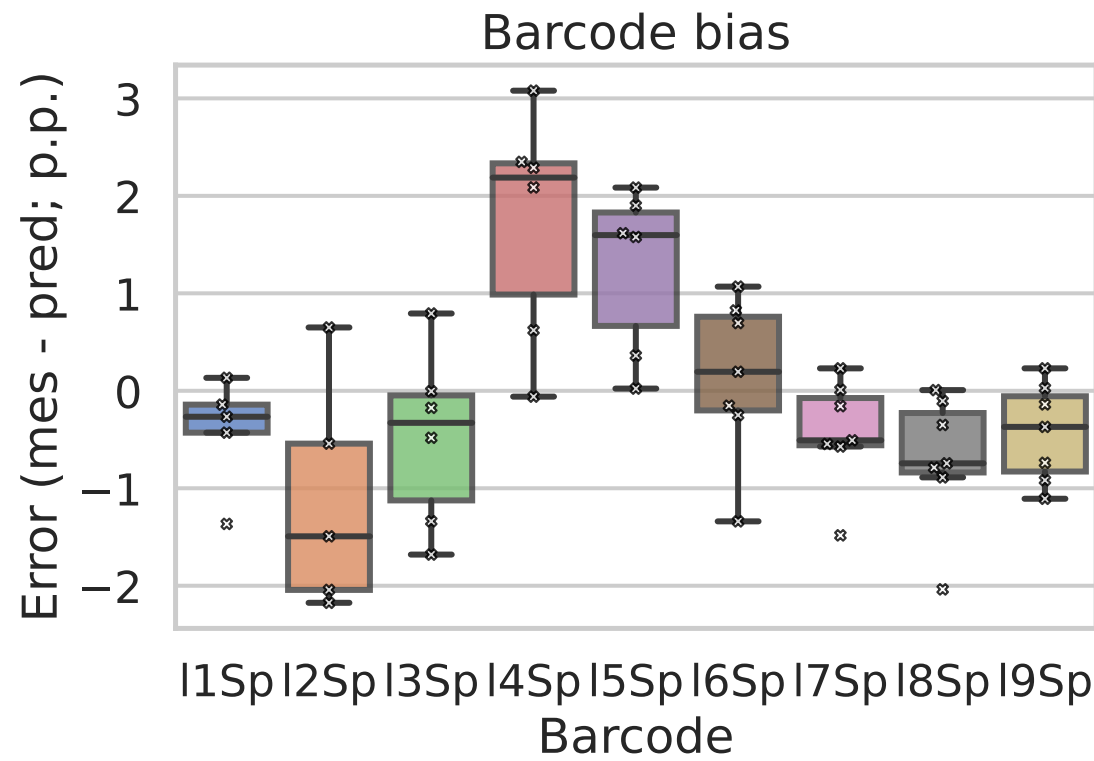
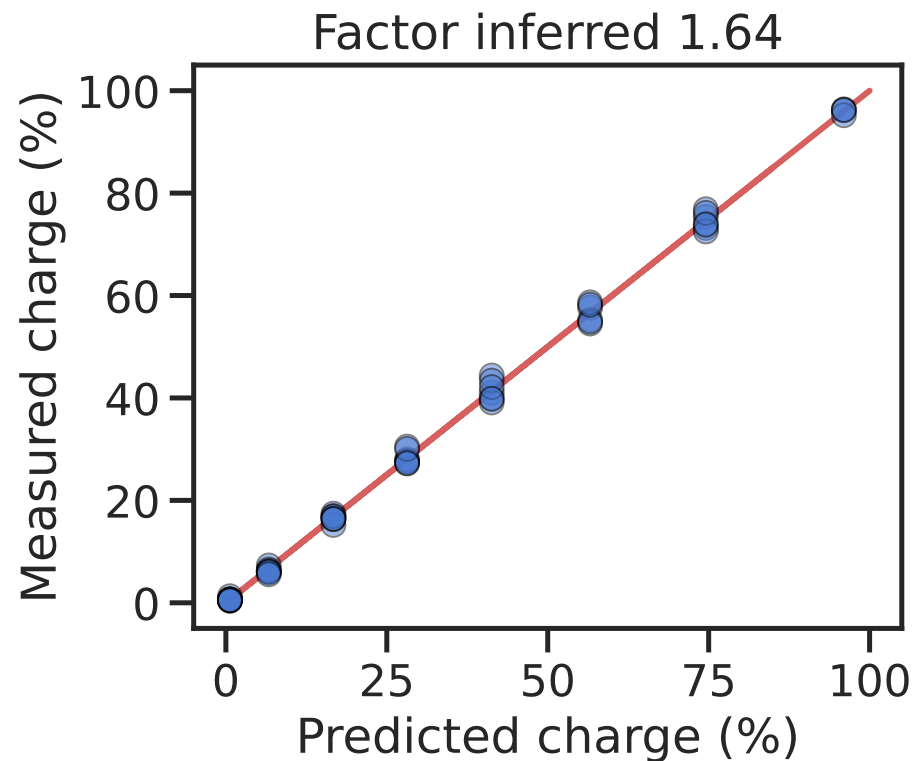
Leu-CAA-2-1



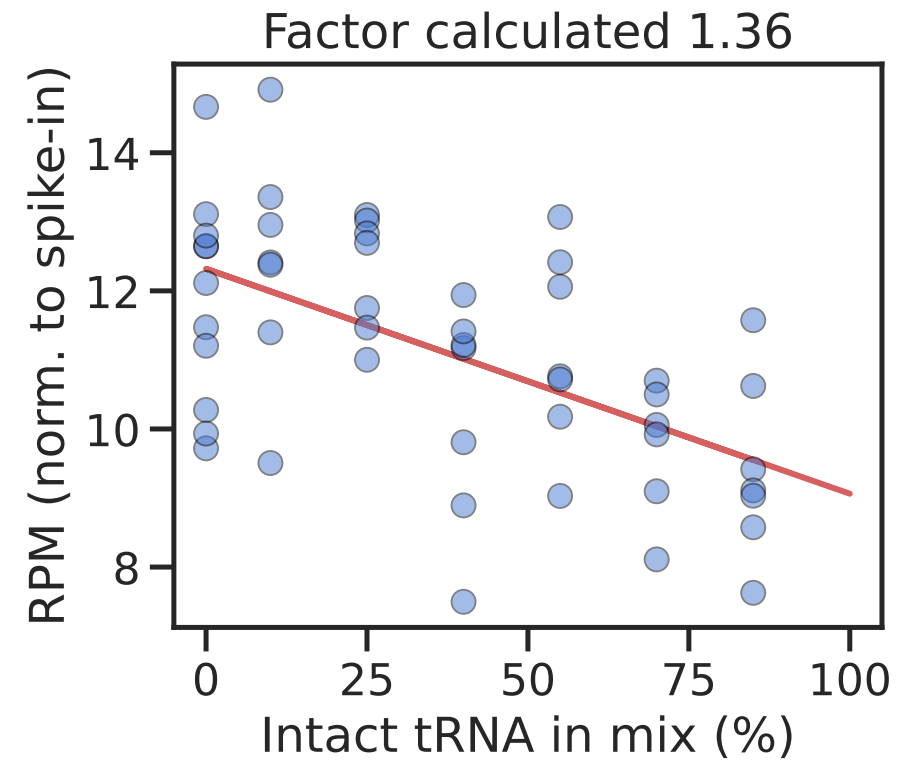
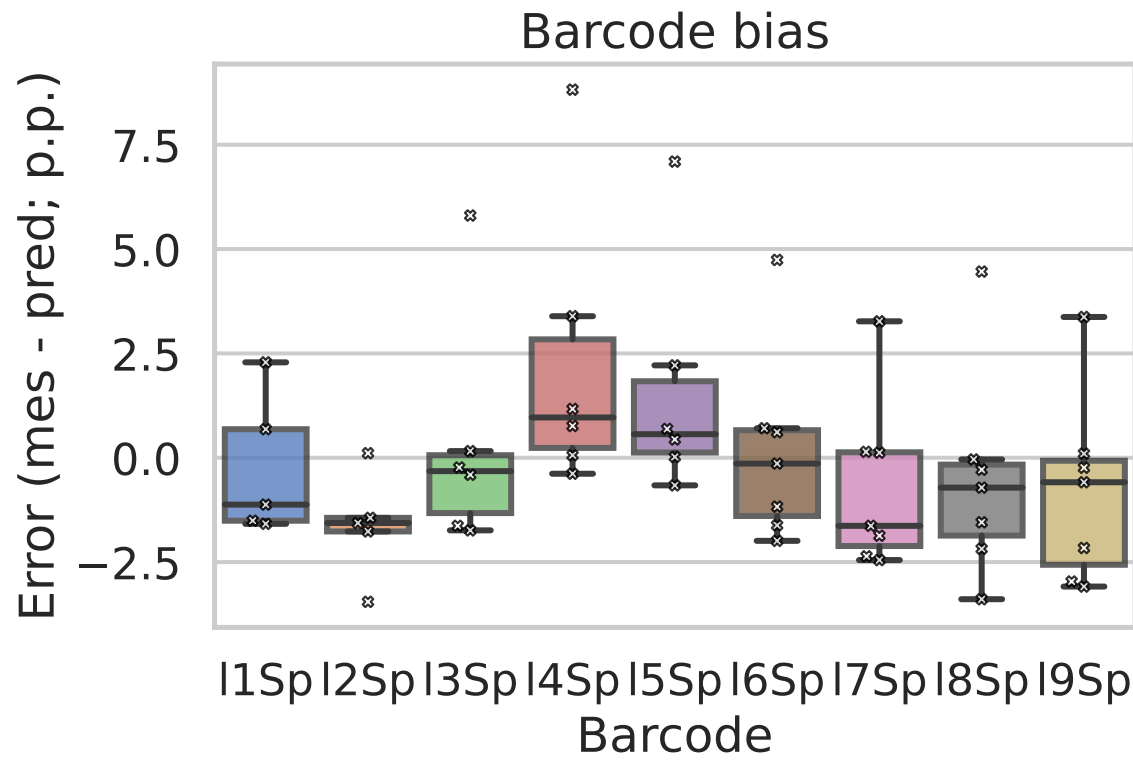
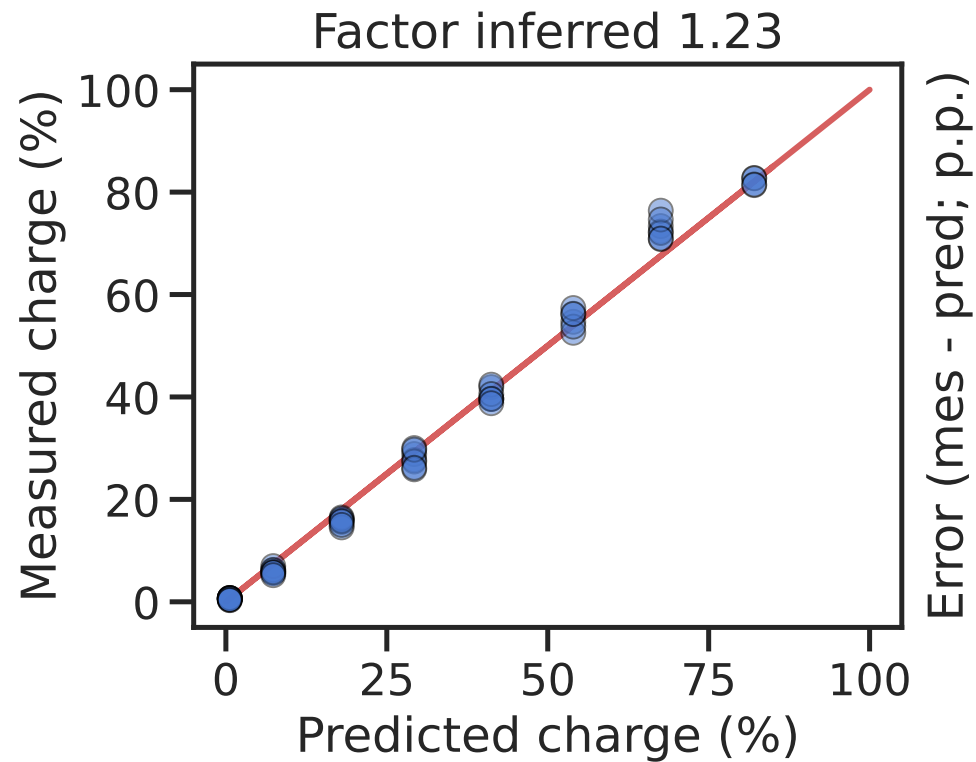
Pro-AGG-2-1



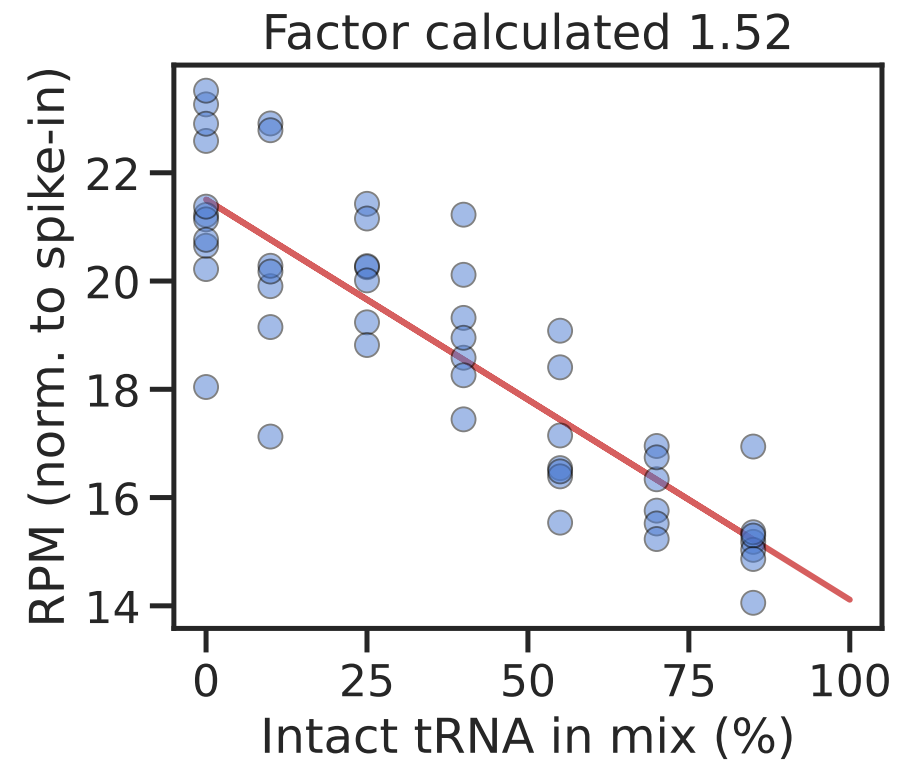
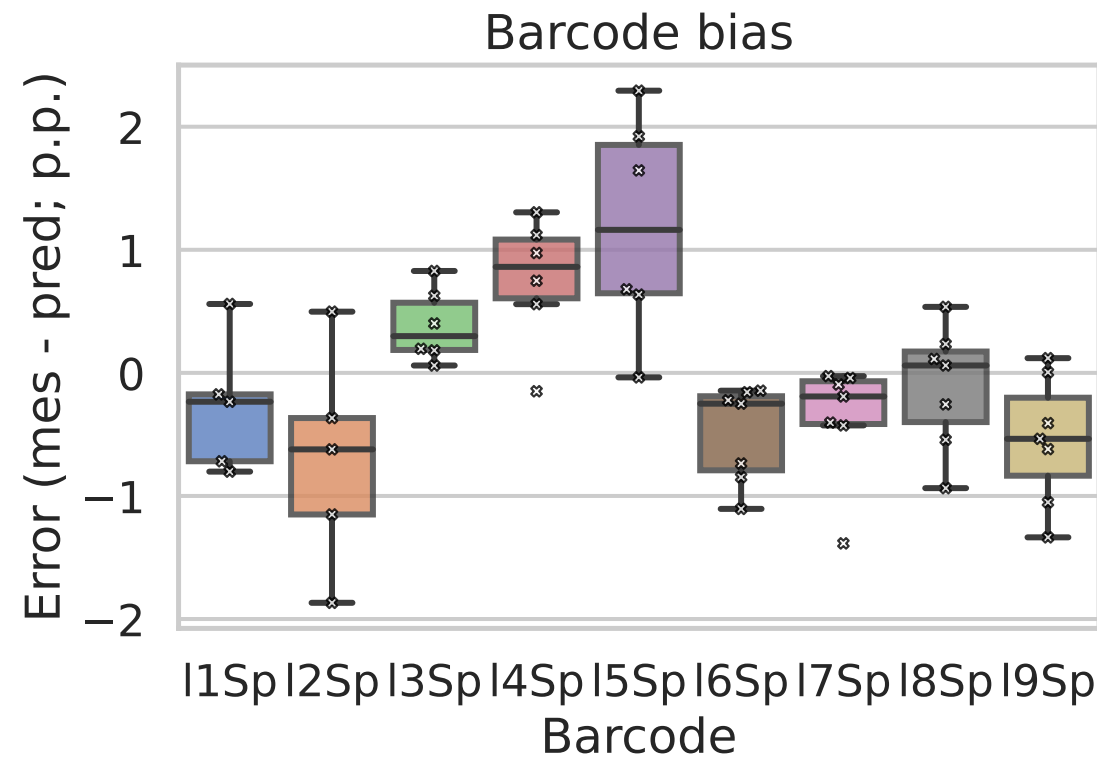
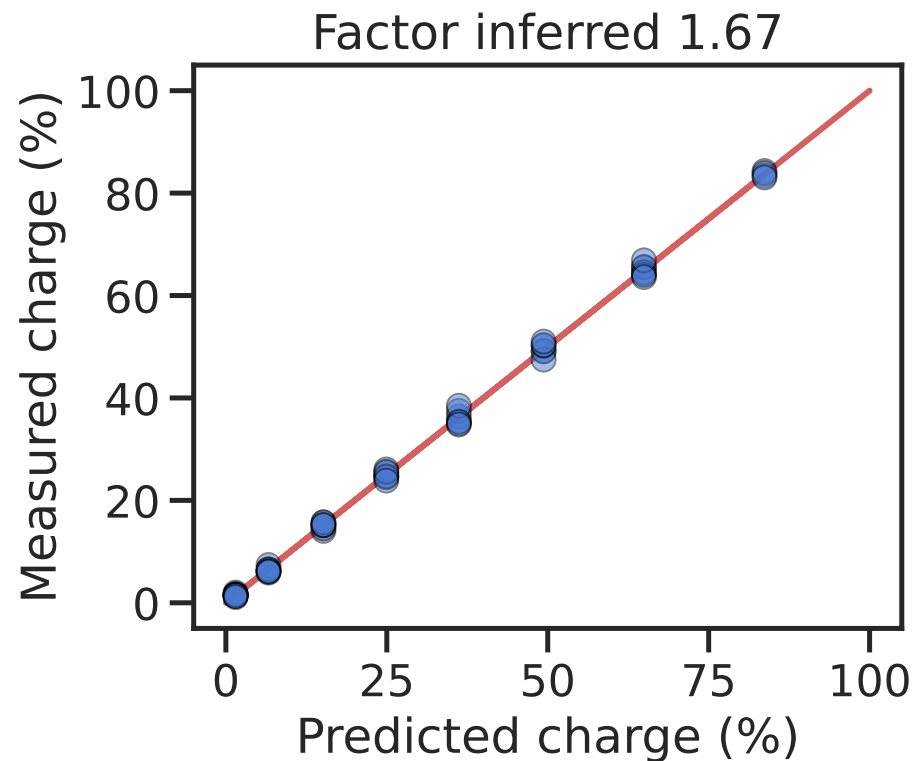
Arg-CCG-2-1



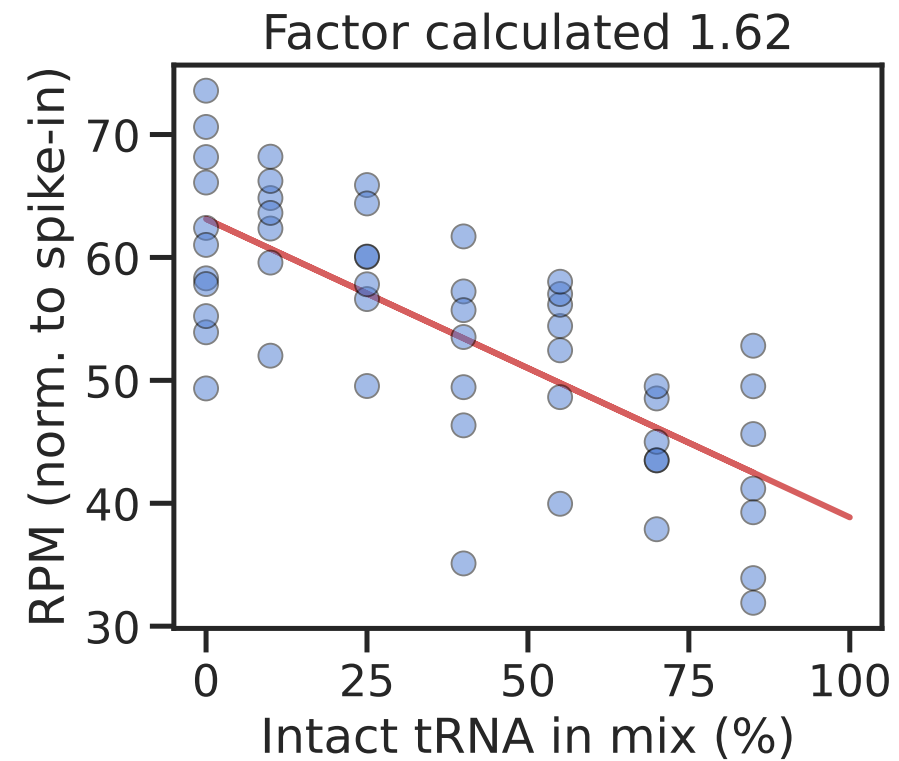
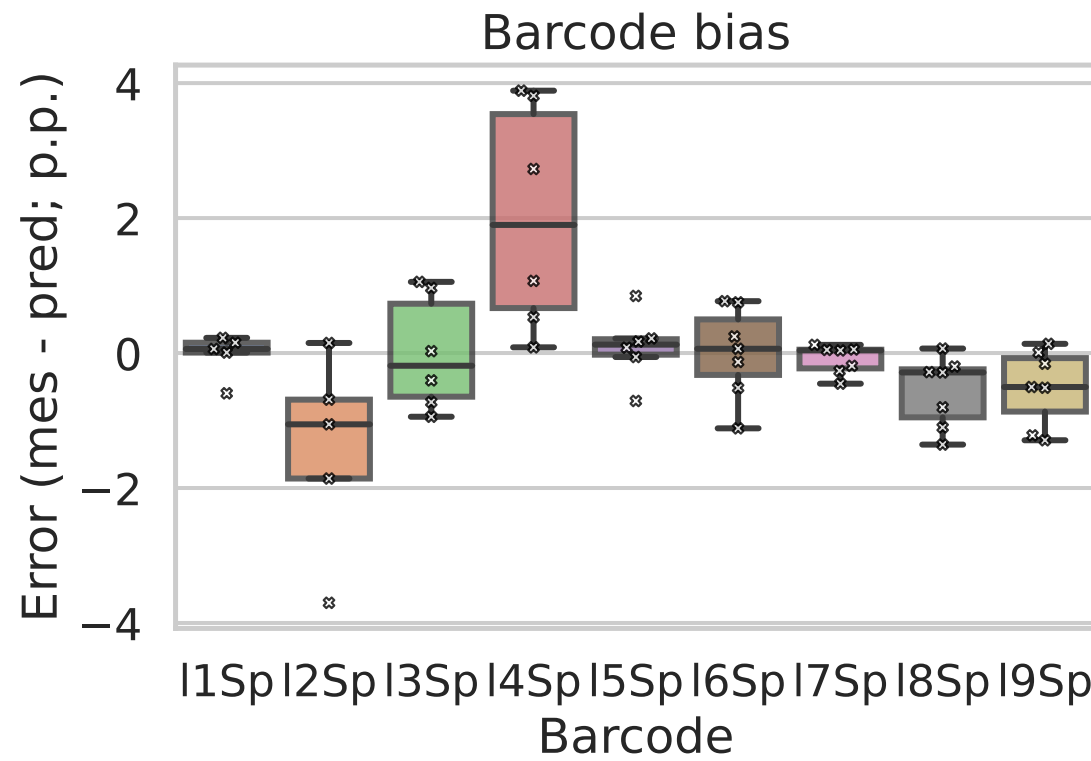
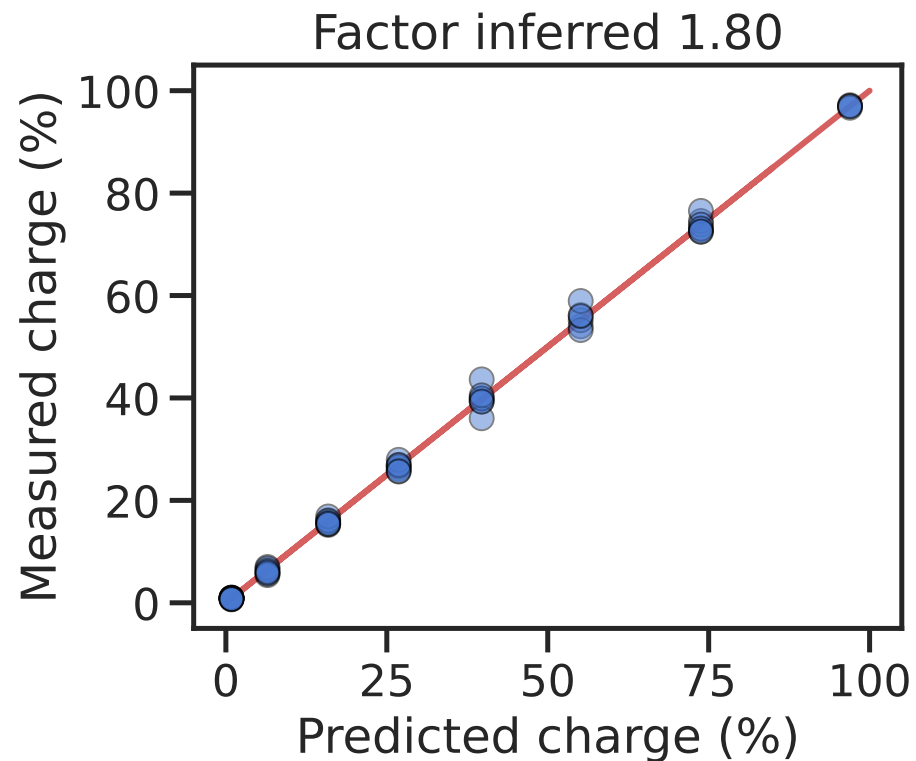
Ser-AGA-2-6

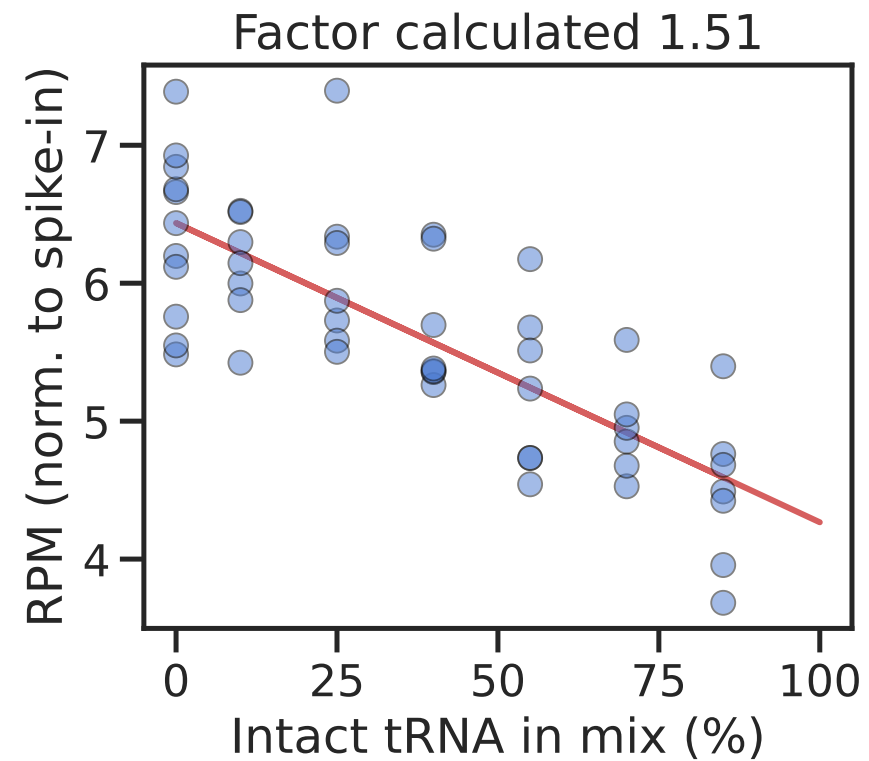
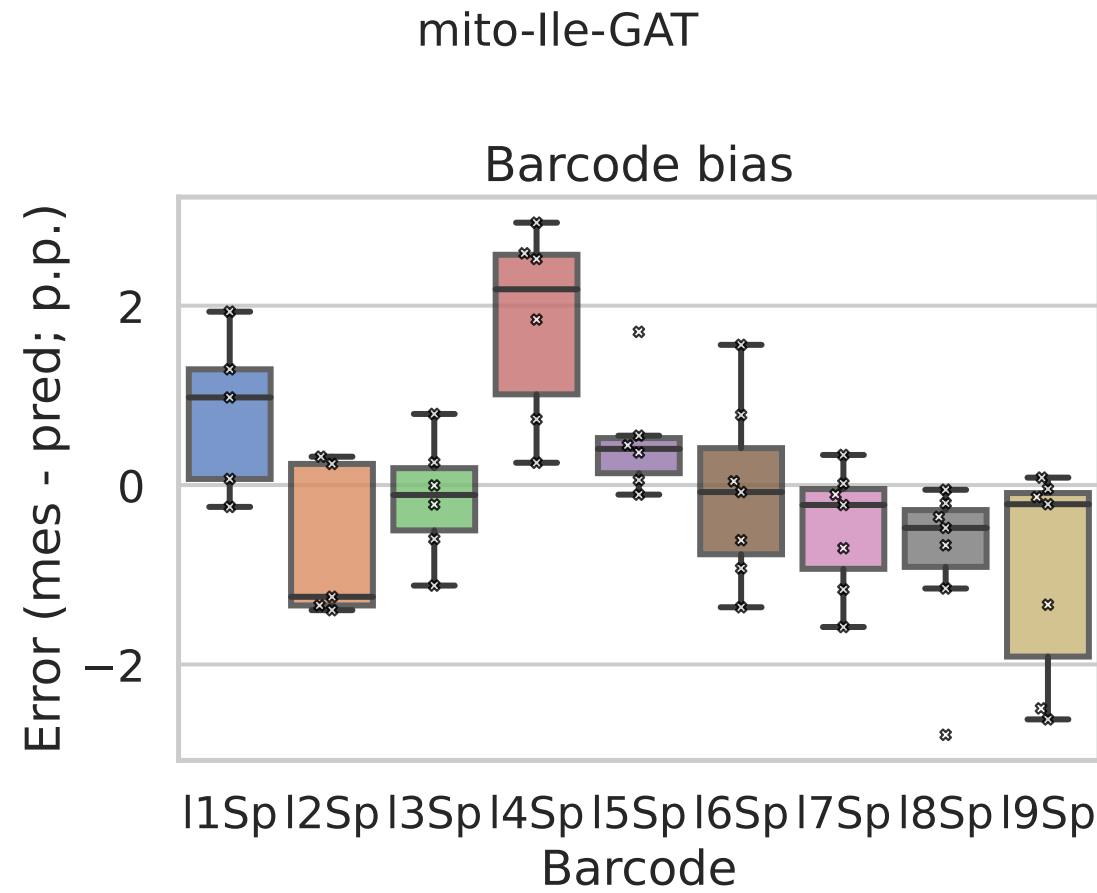
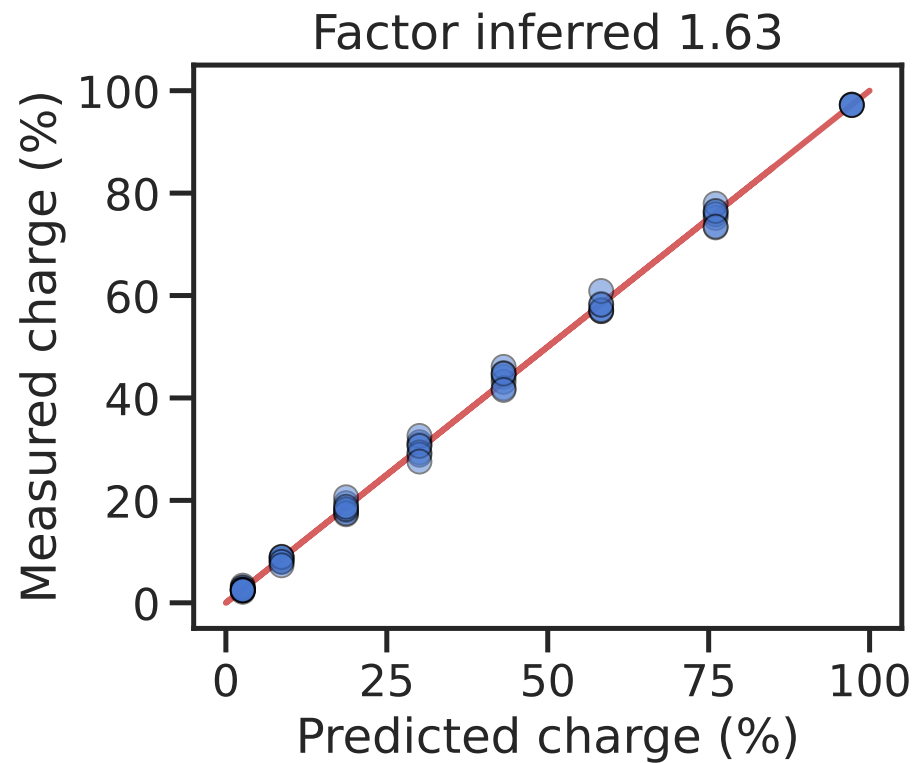


mito-Val-TAC

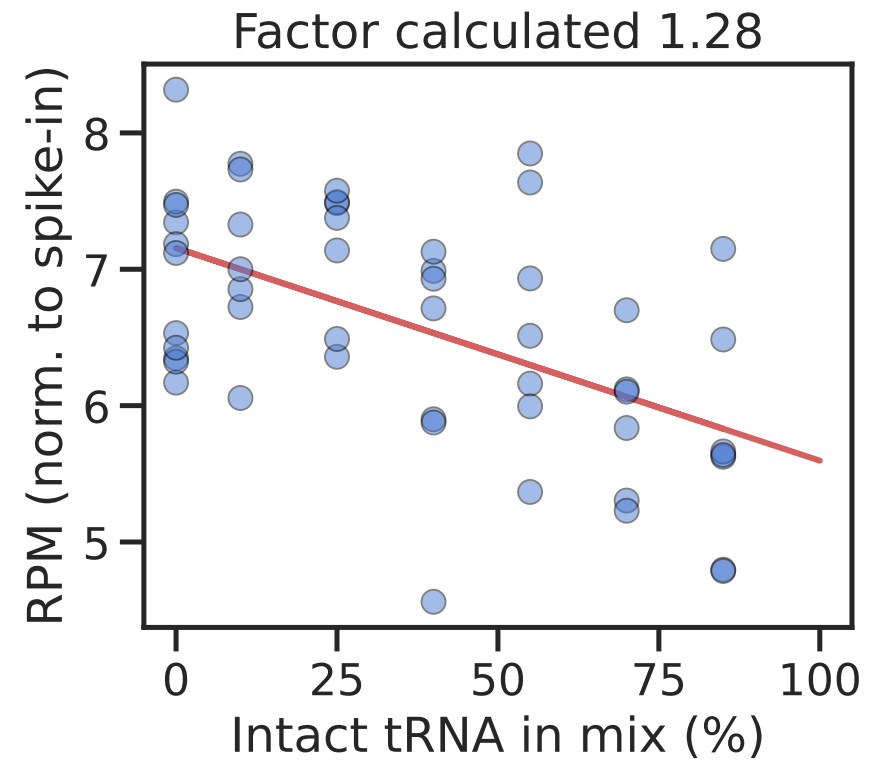
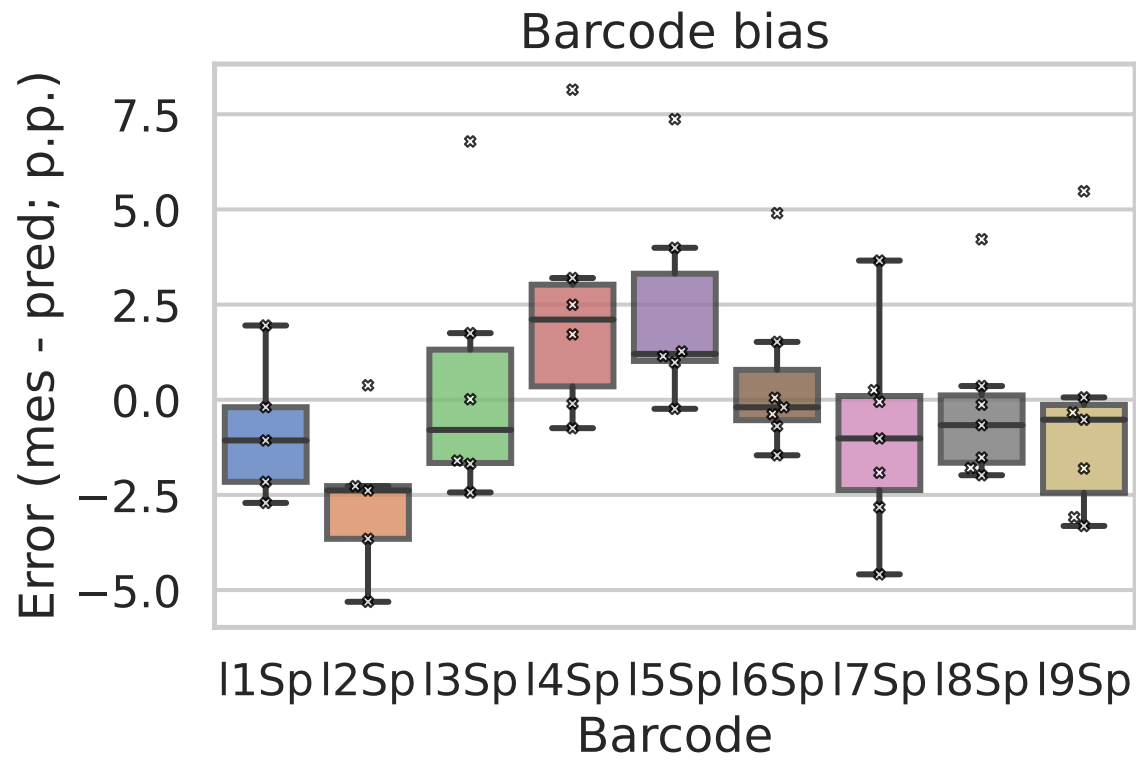
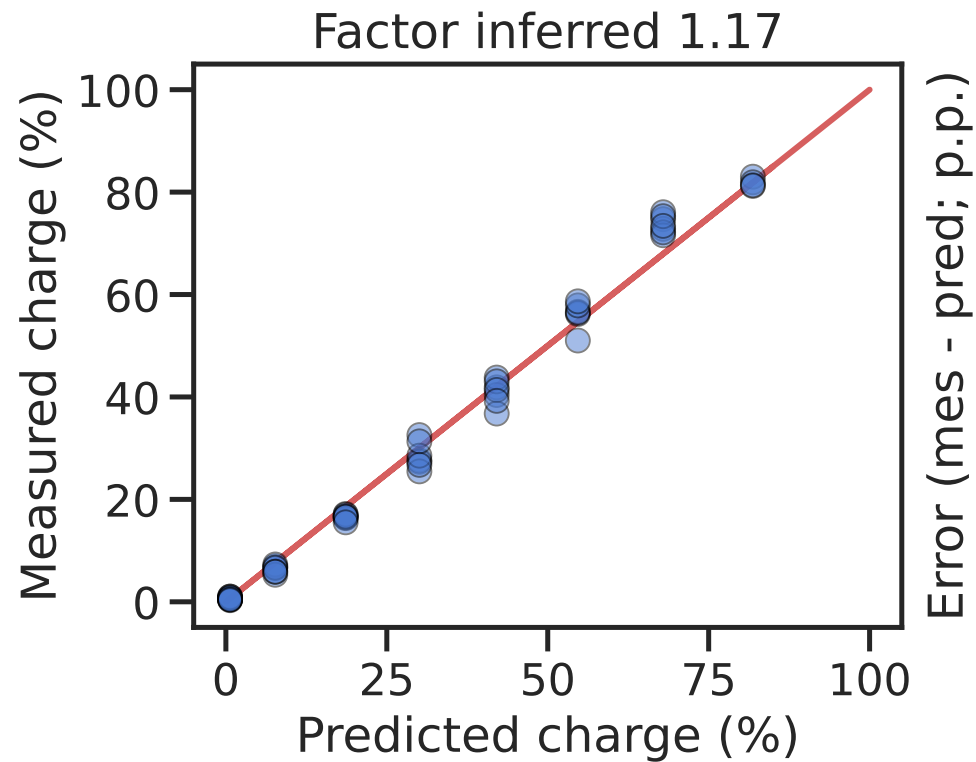


Val-CAC-1-8

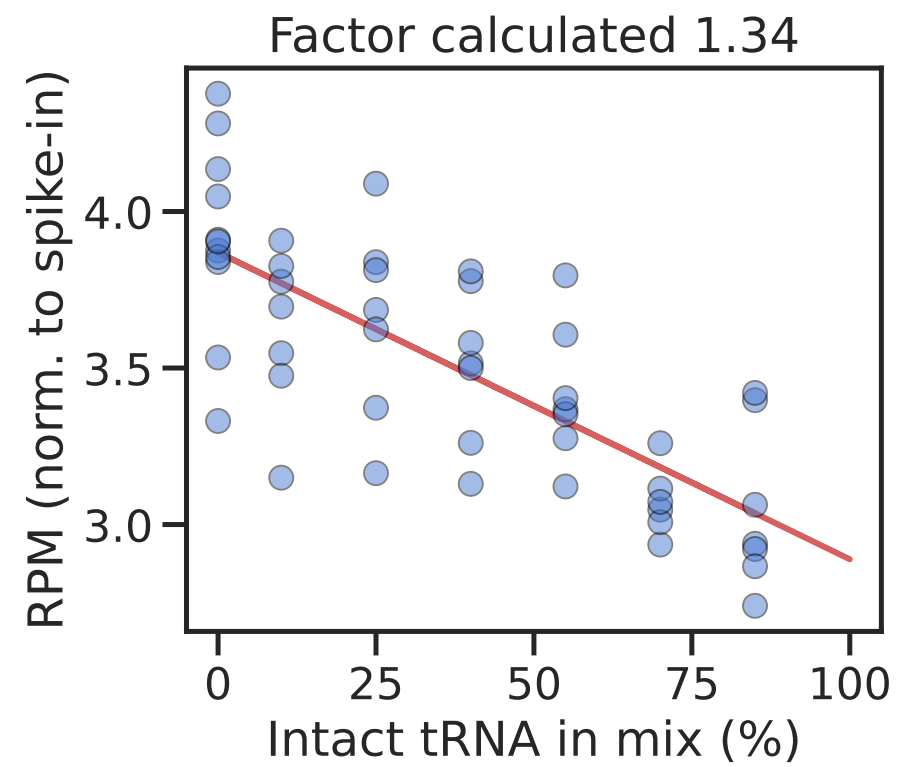
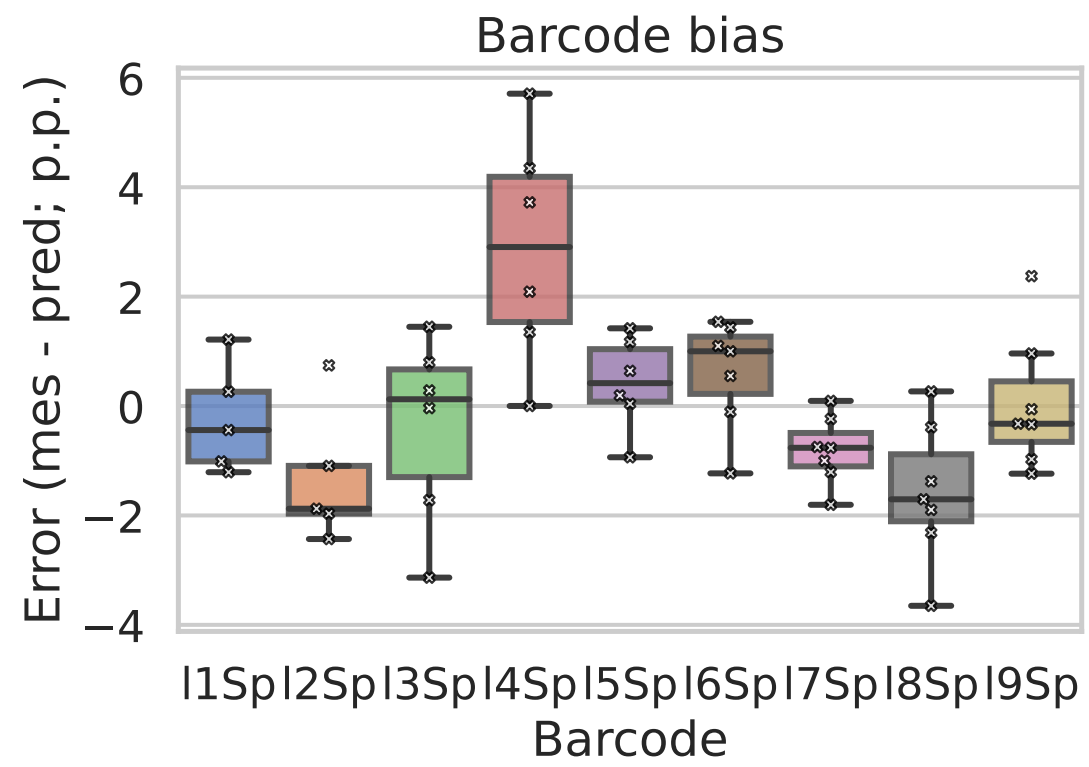
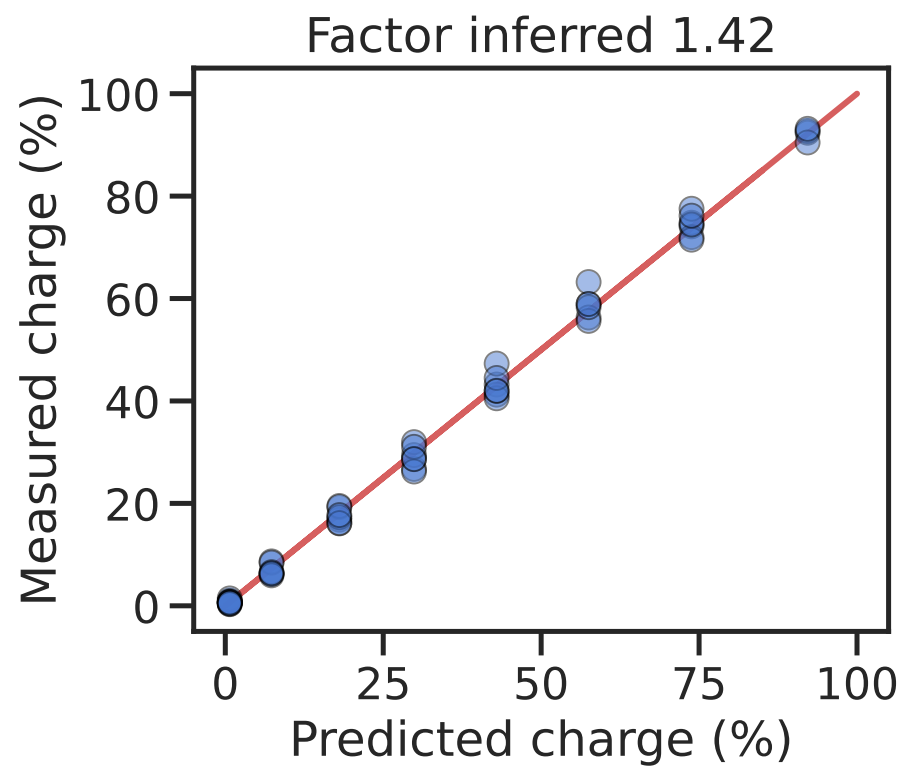




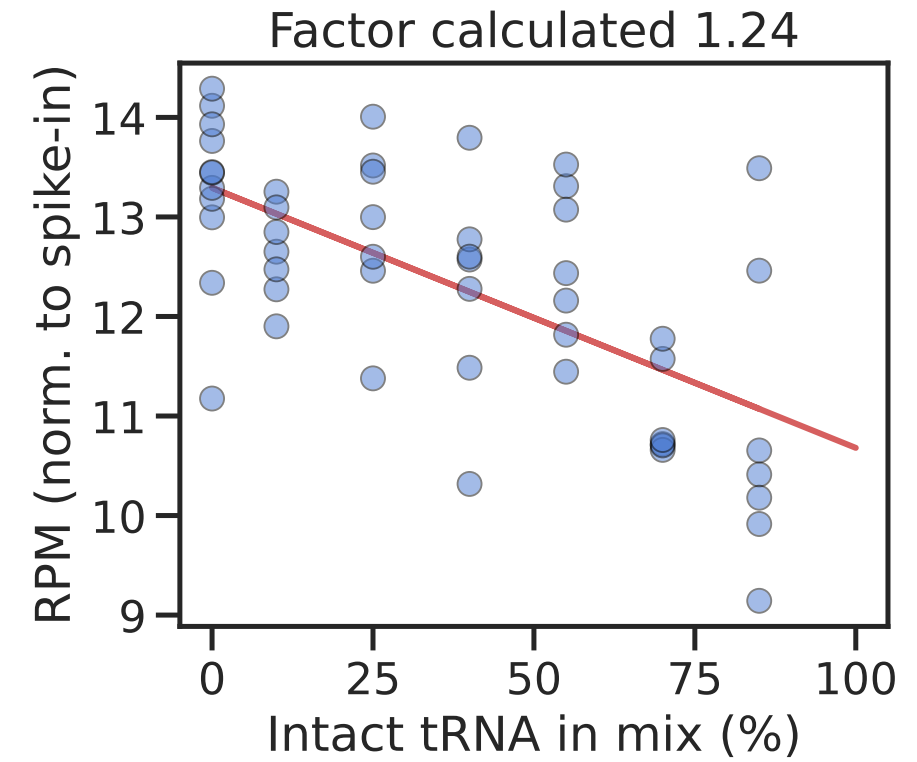
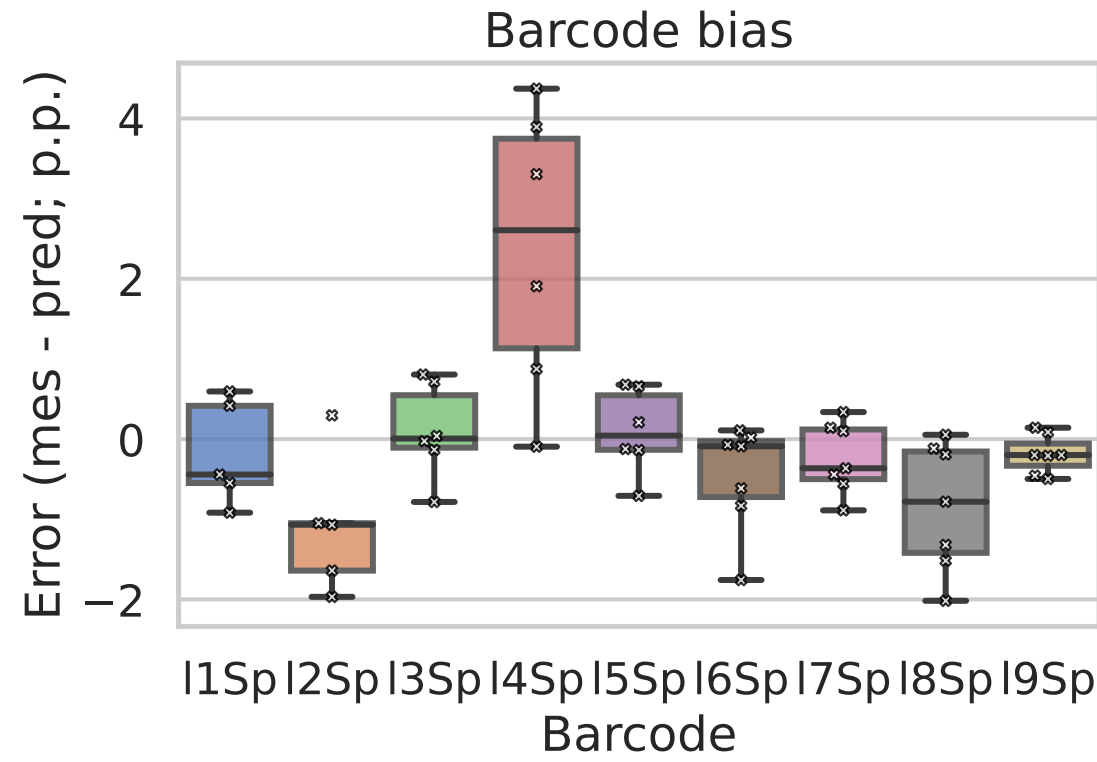
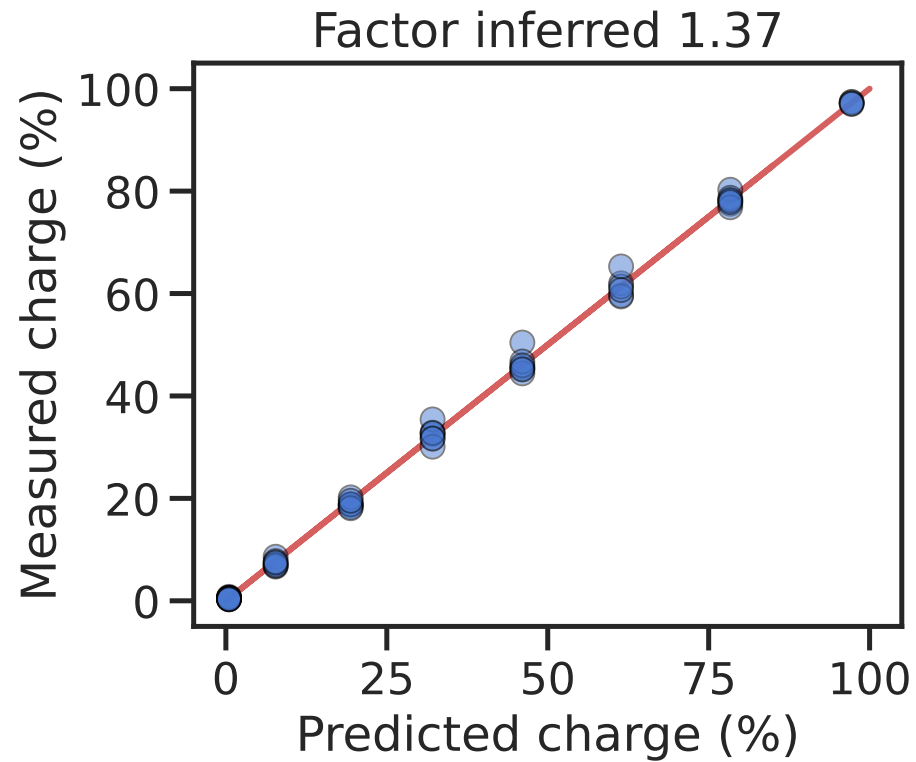
Ser-AGA-1-1



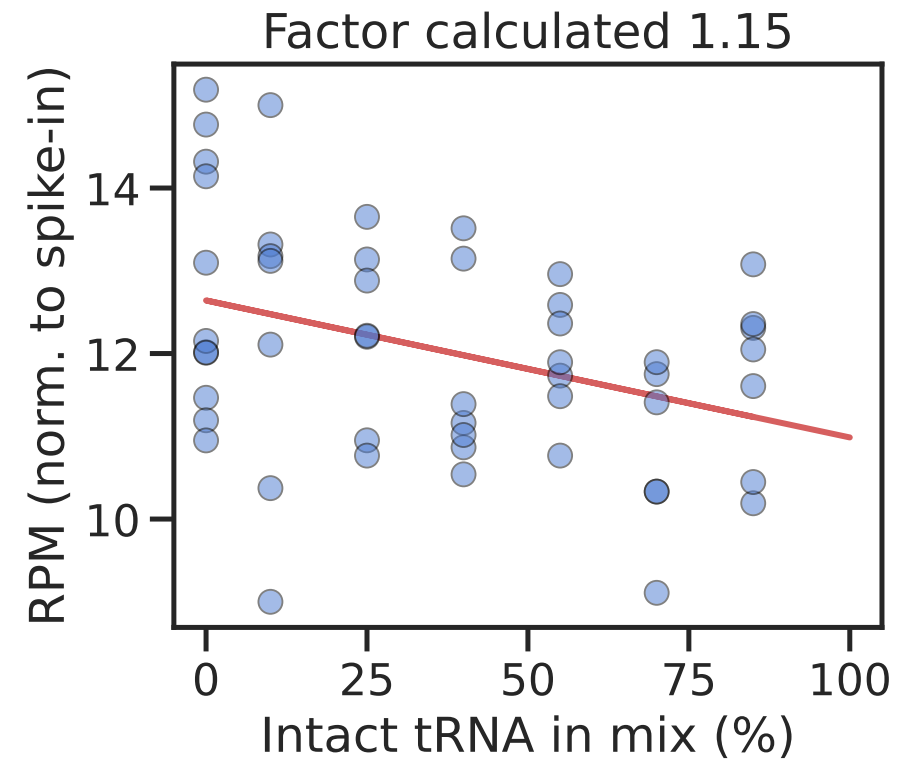
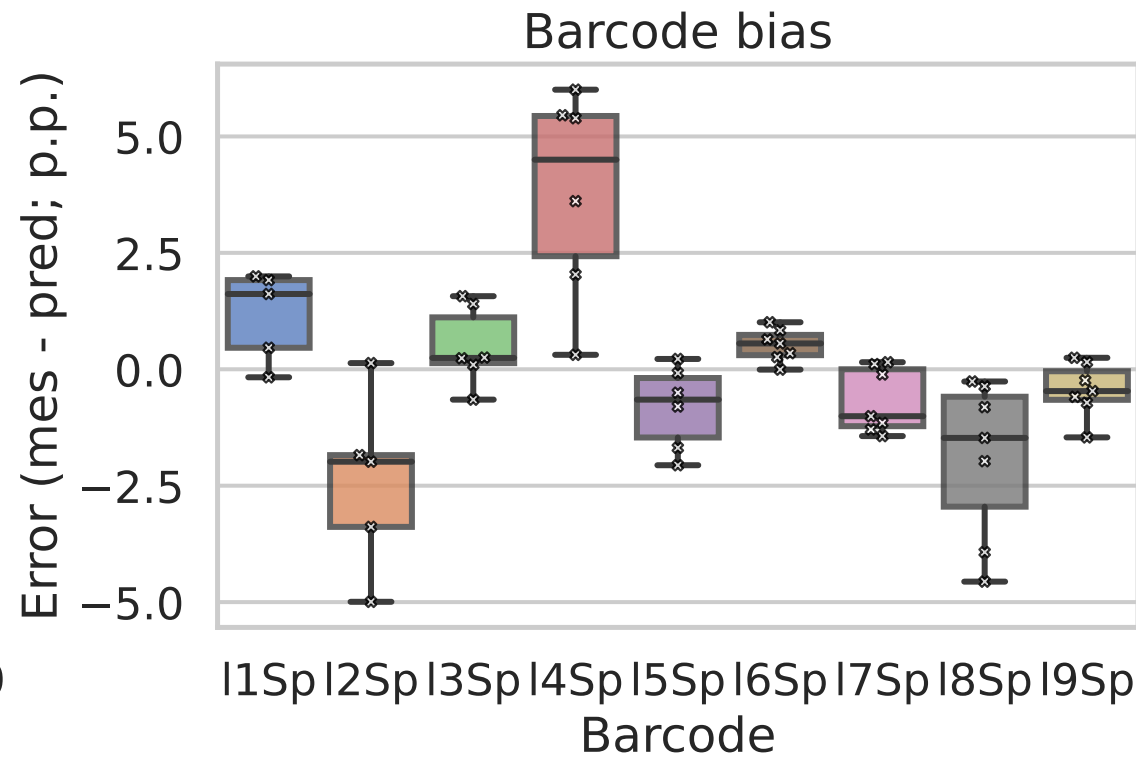
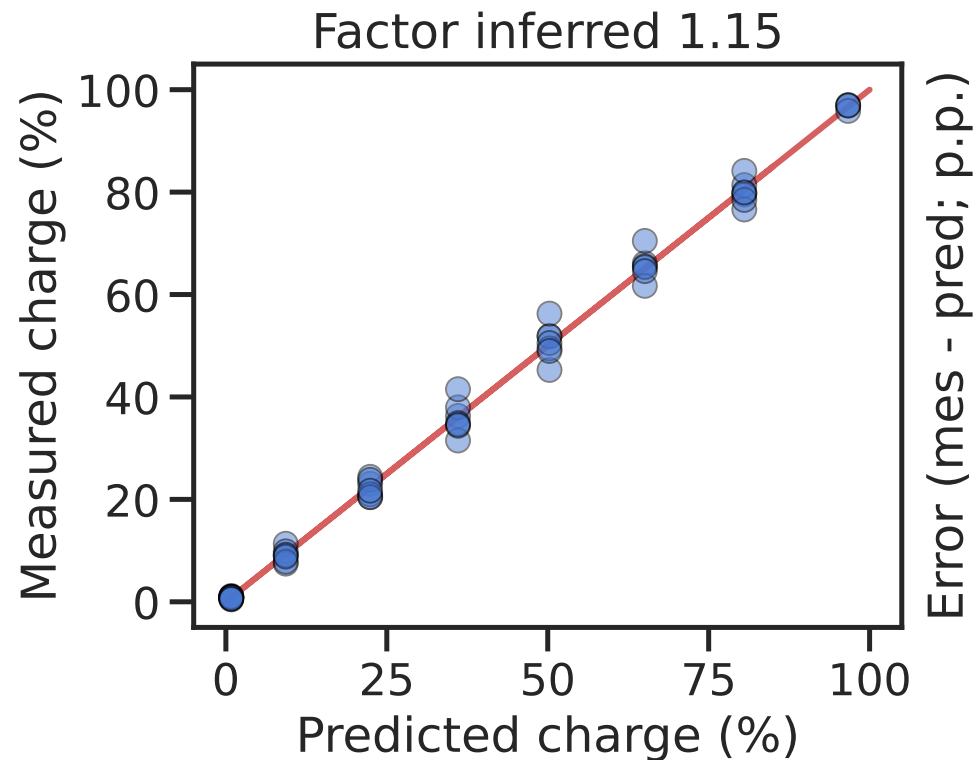
Thr-AGT-4-1



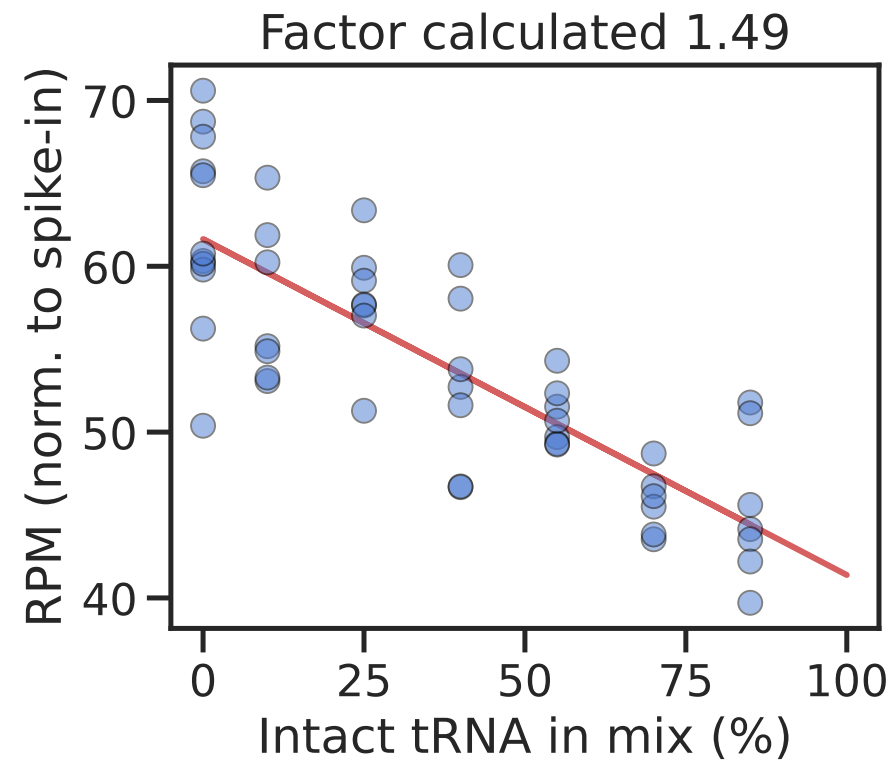
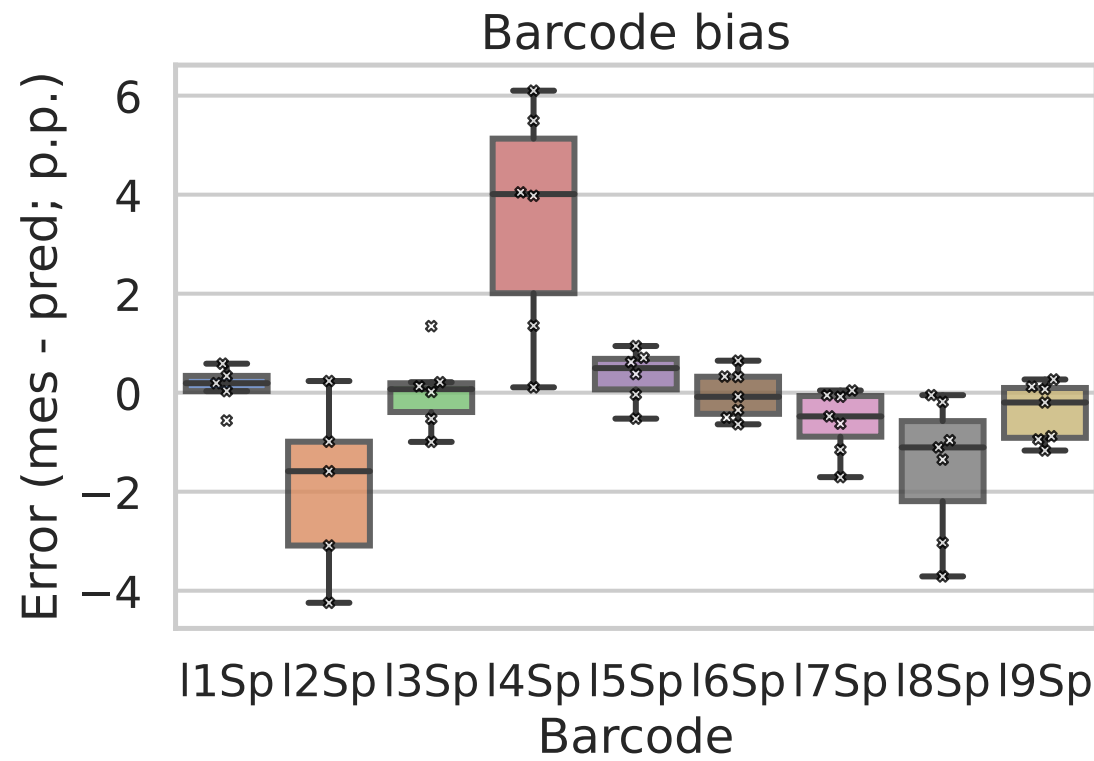
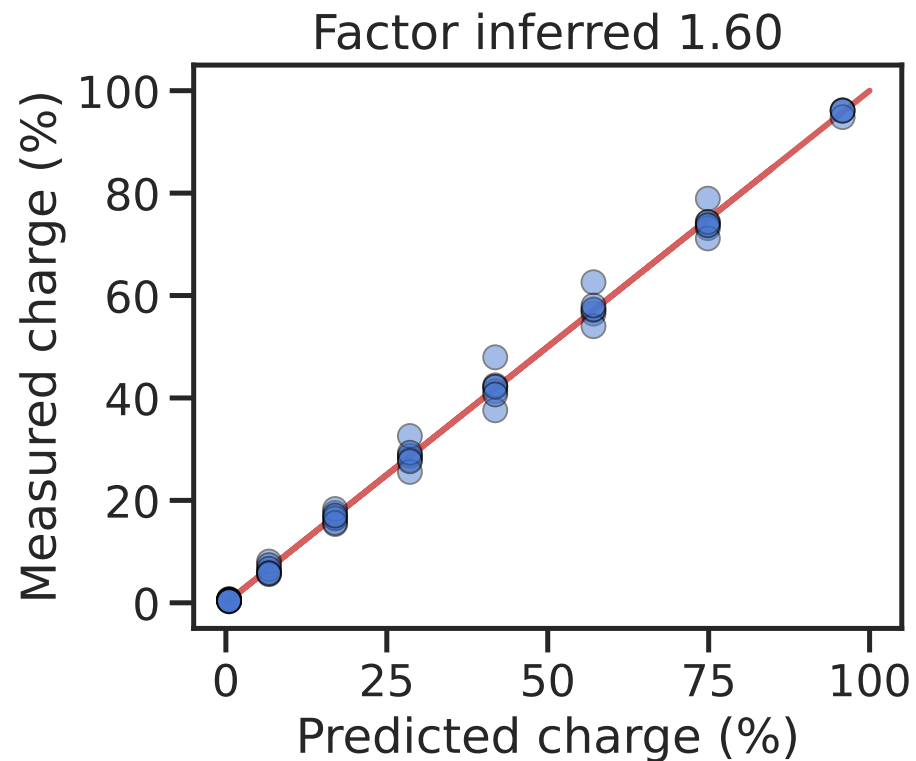
Ala-TGC-5-1



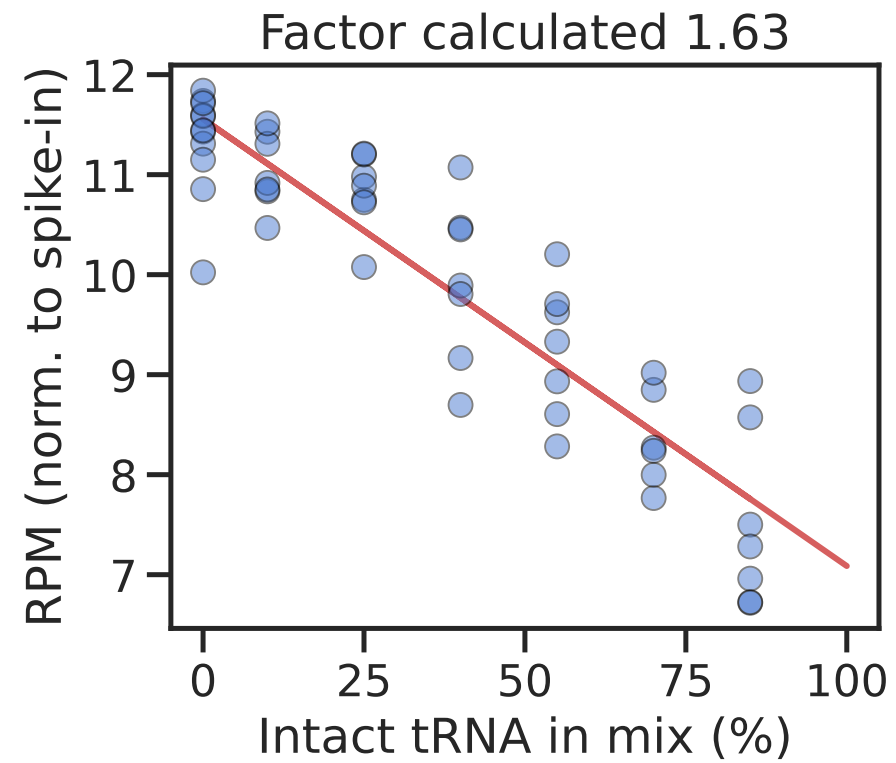
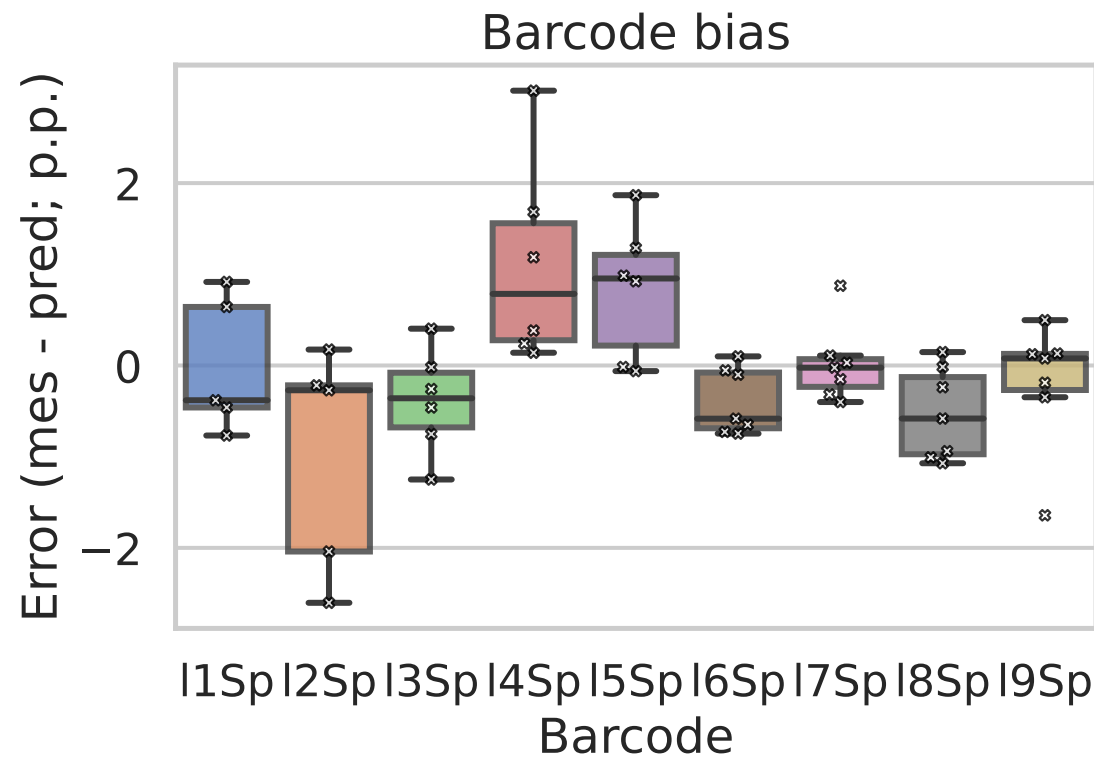
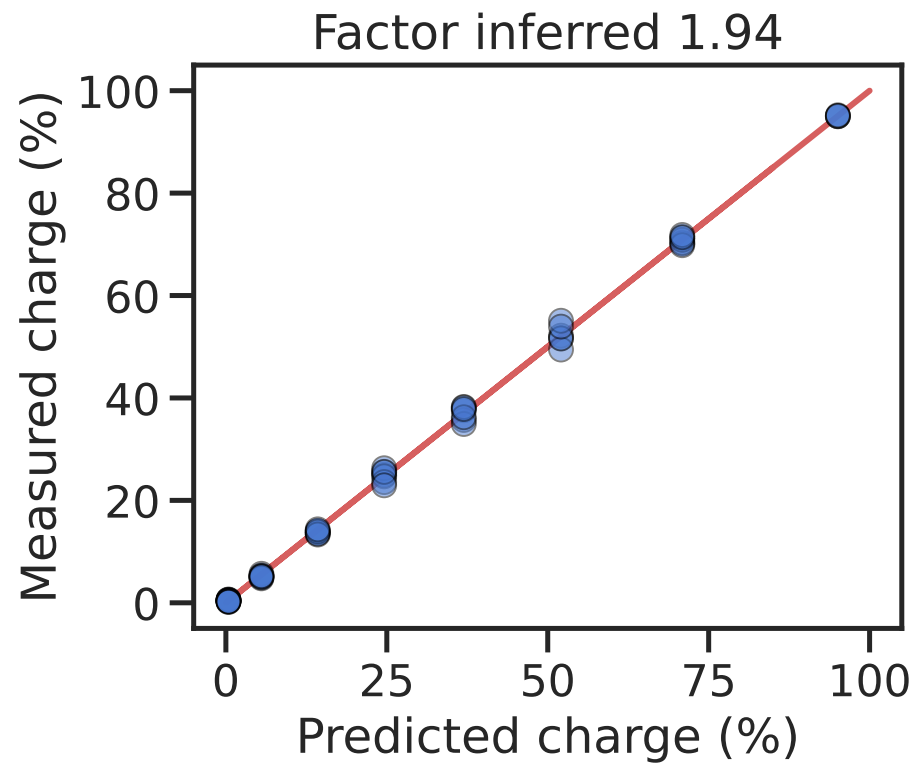
Cys-GCA-2-2



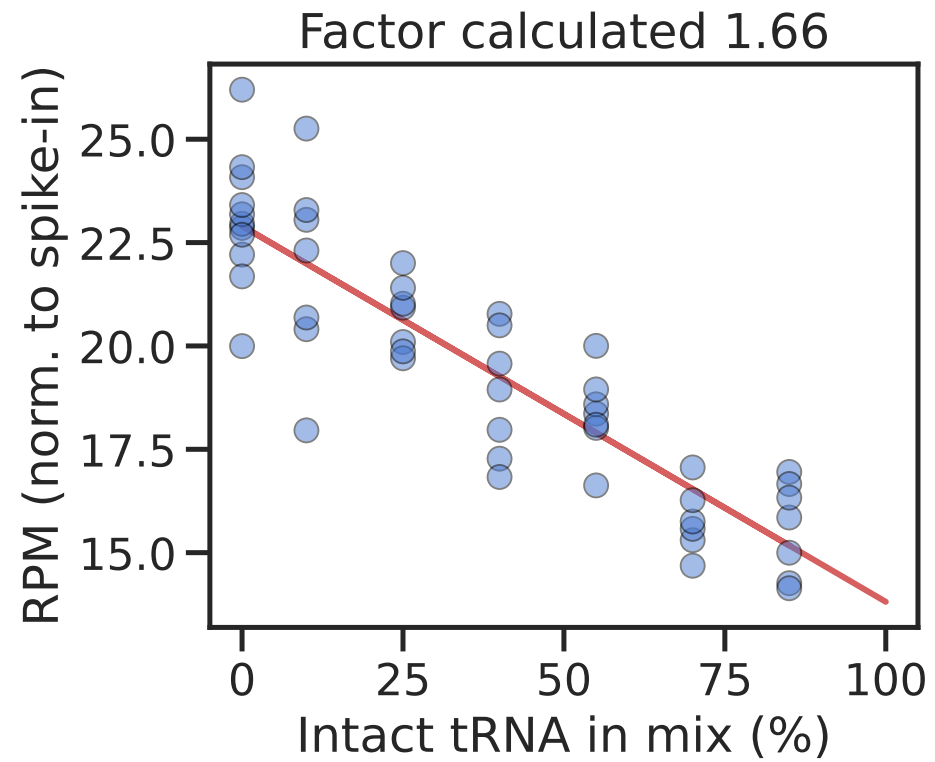
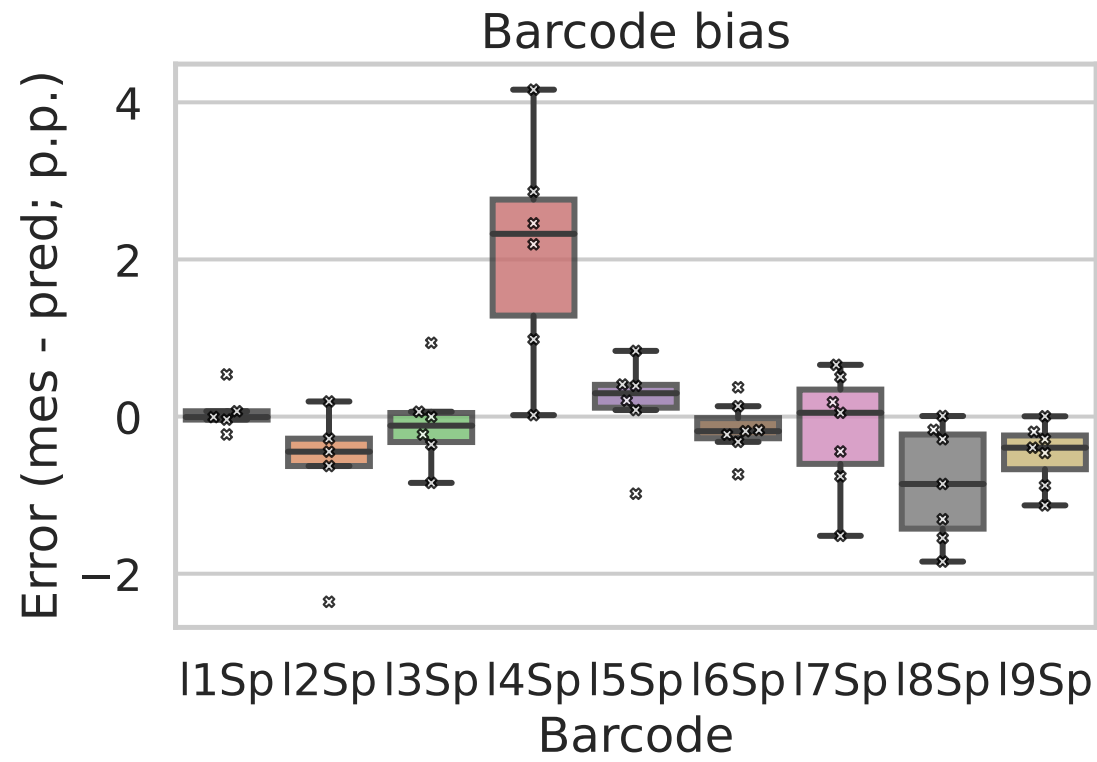
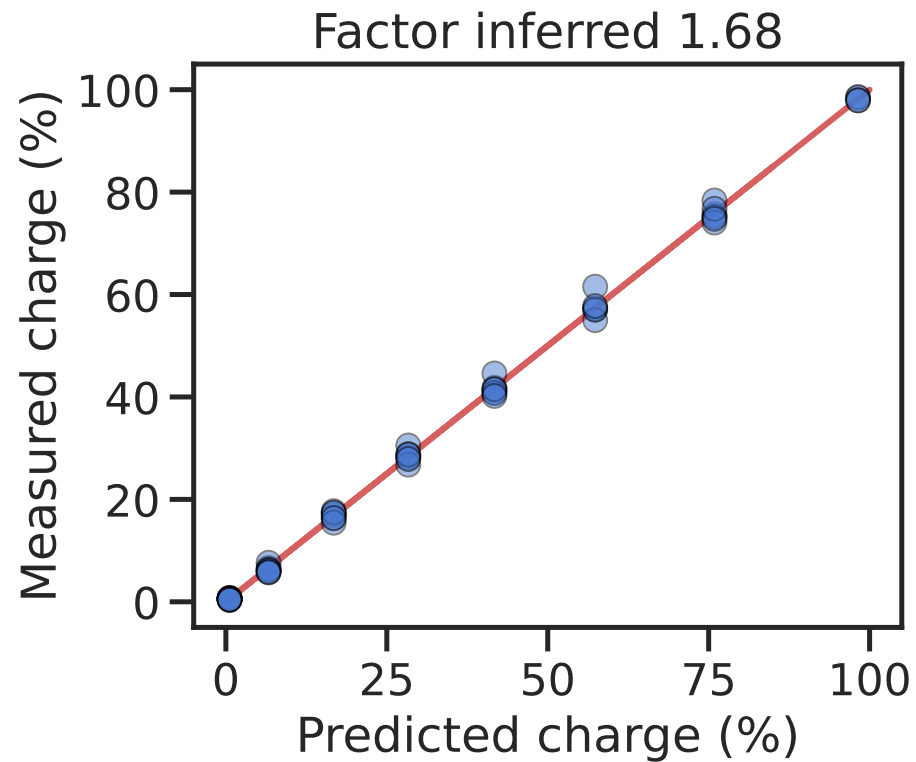
Gln-CTG-1-4



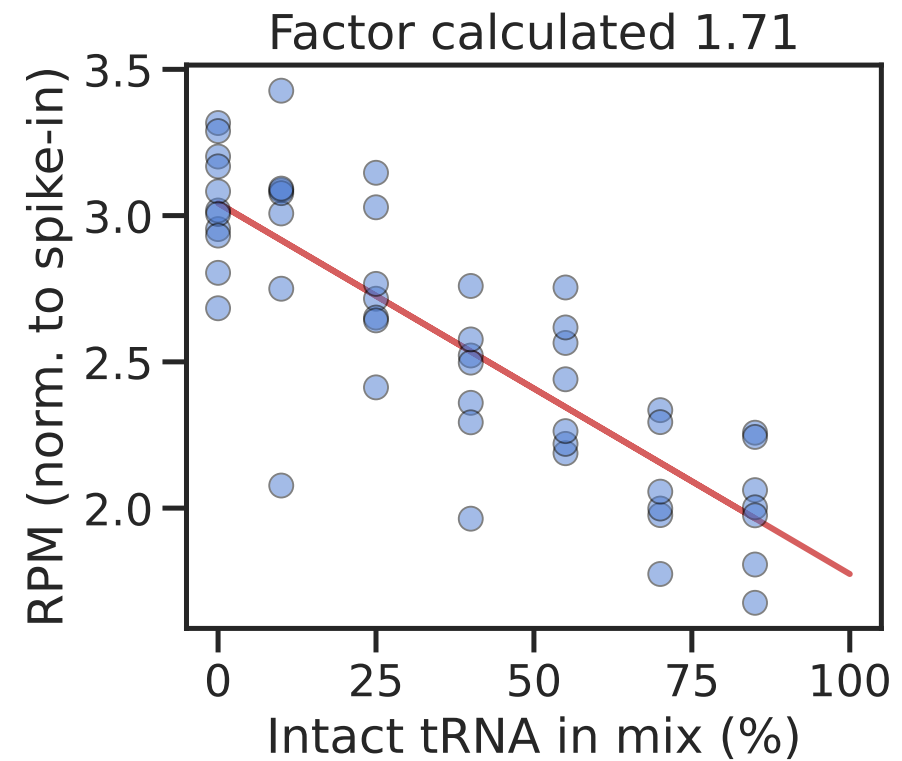
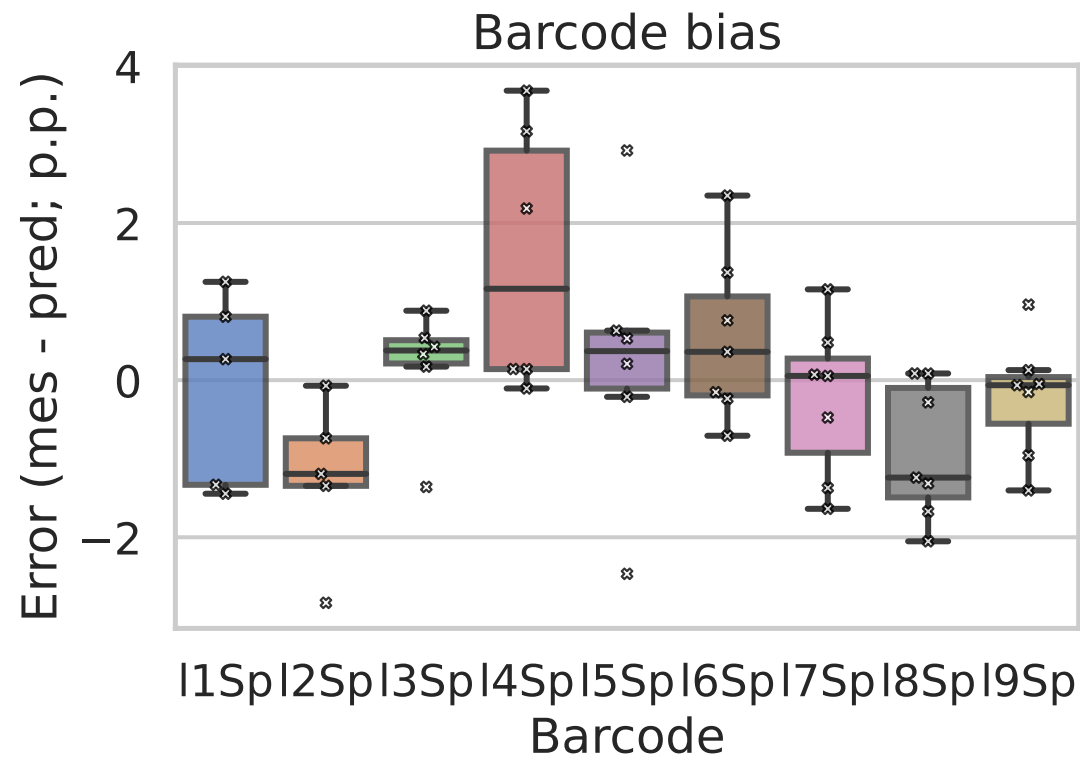
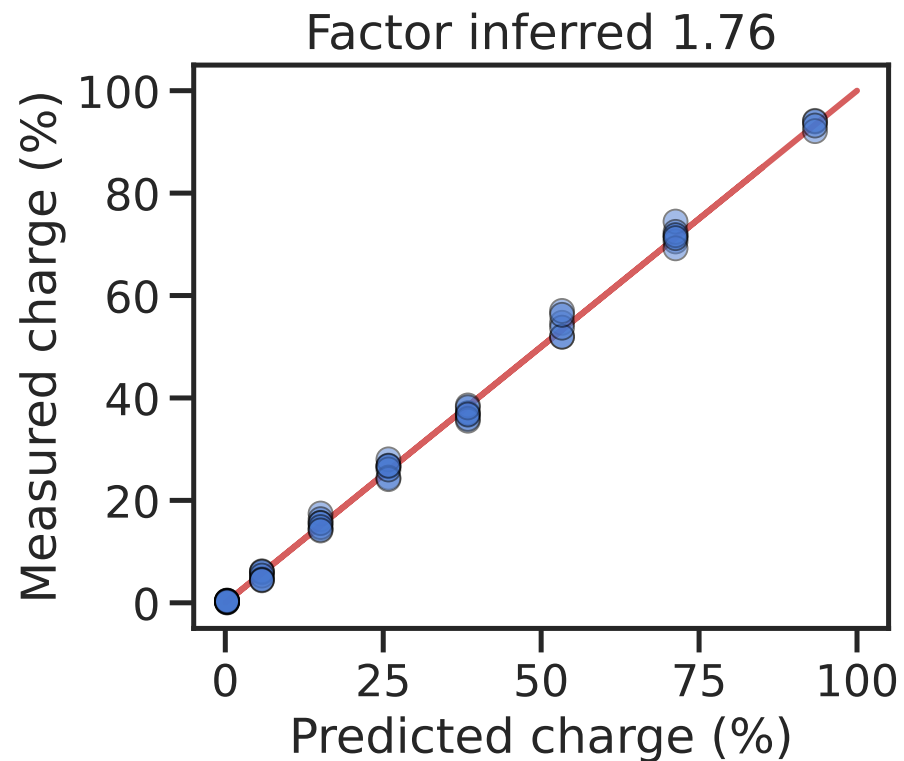
Arg-TCG-2-1



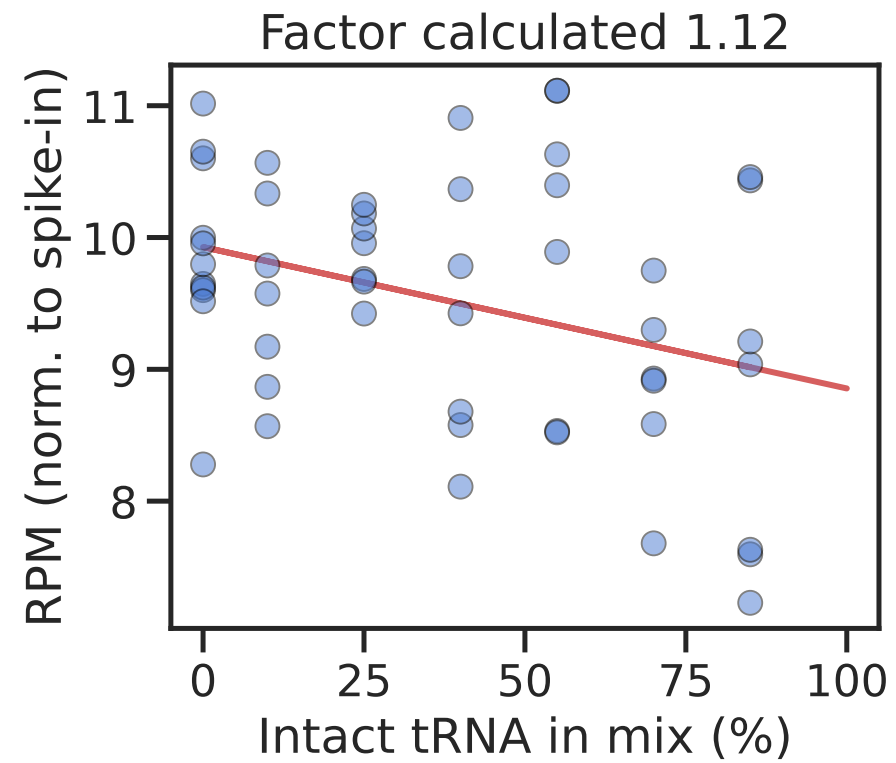
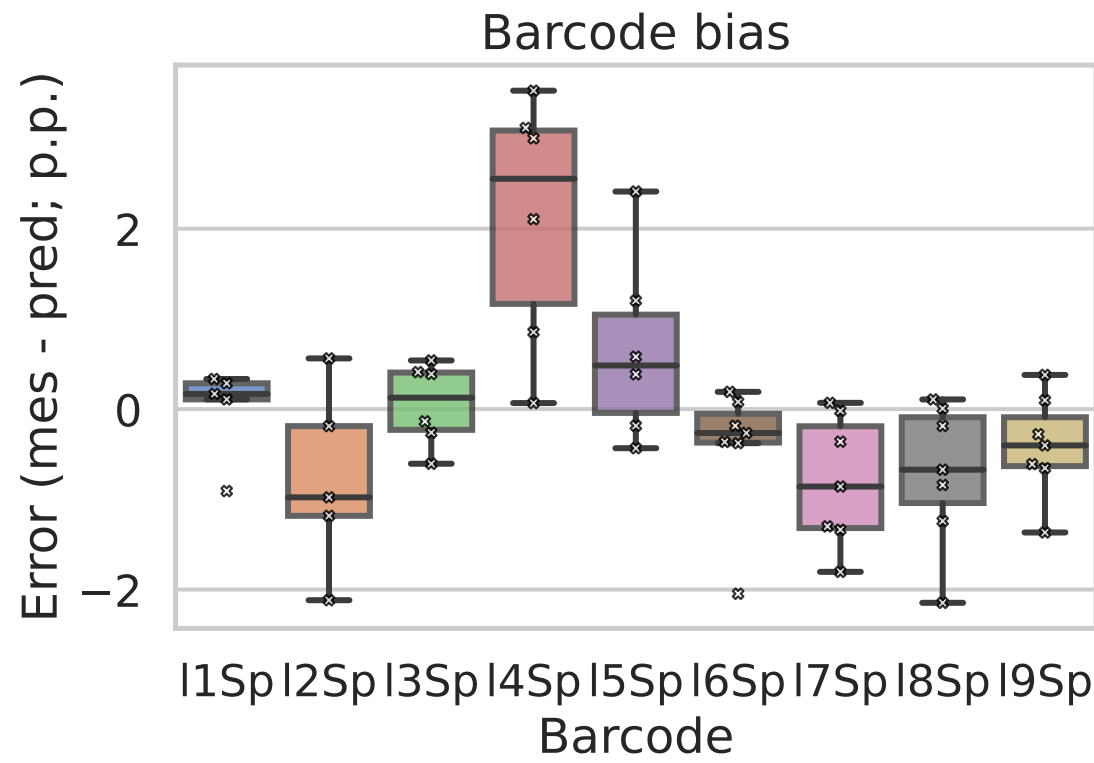
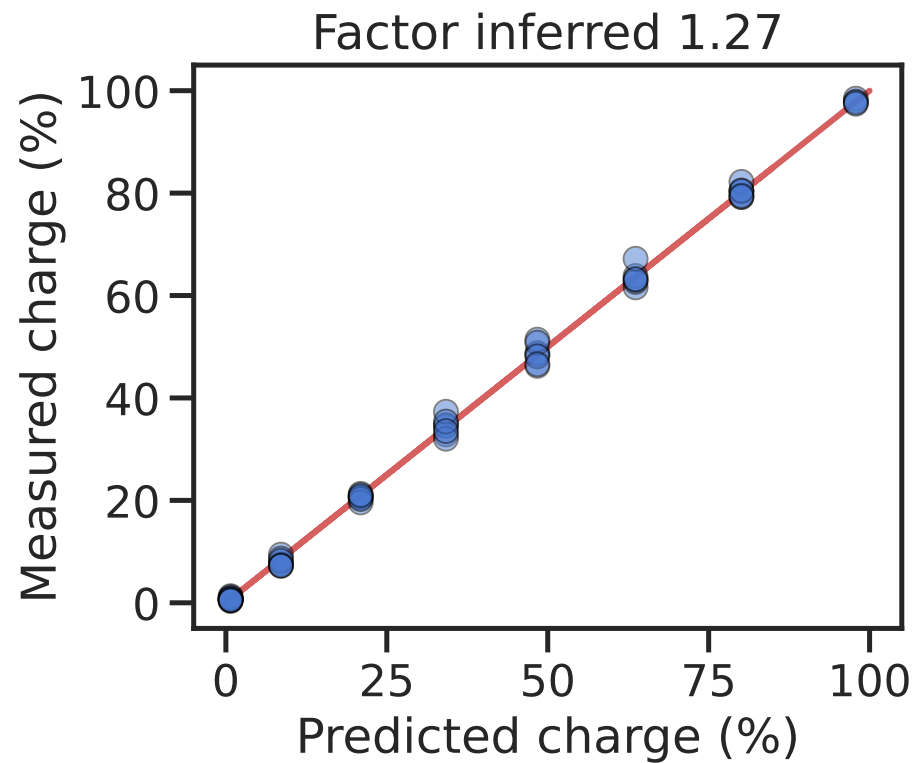
Gly-CCC-1-1



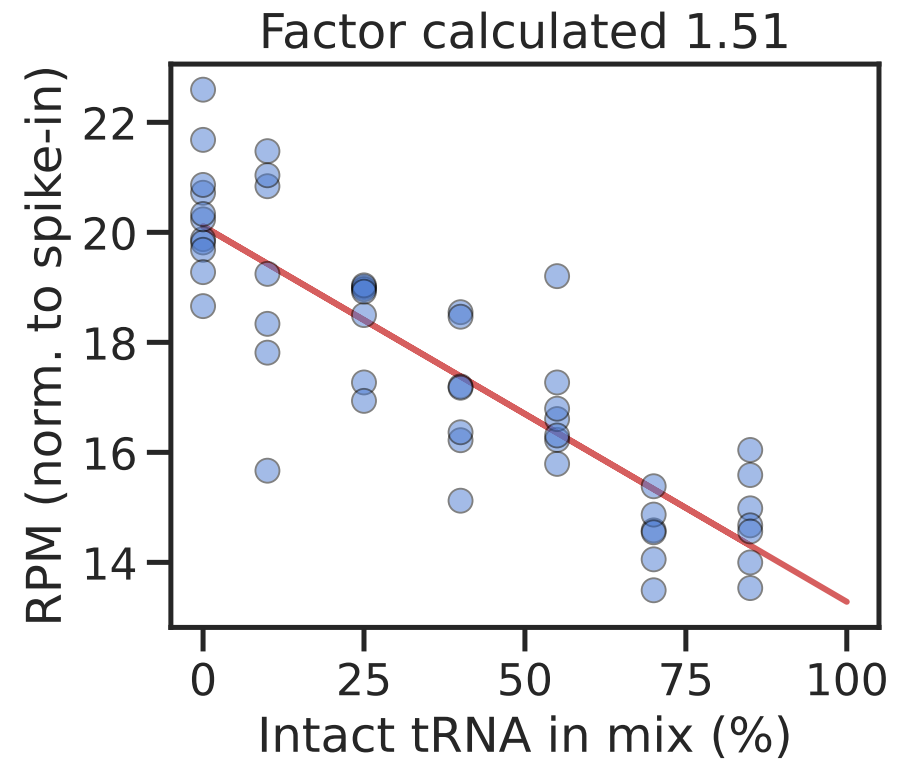
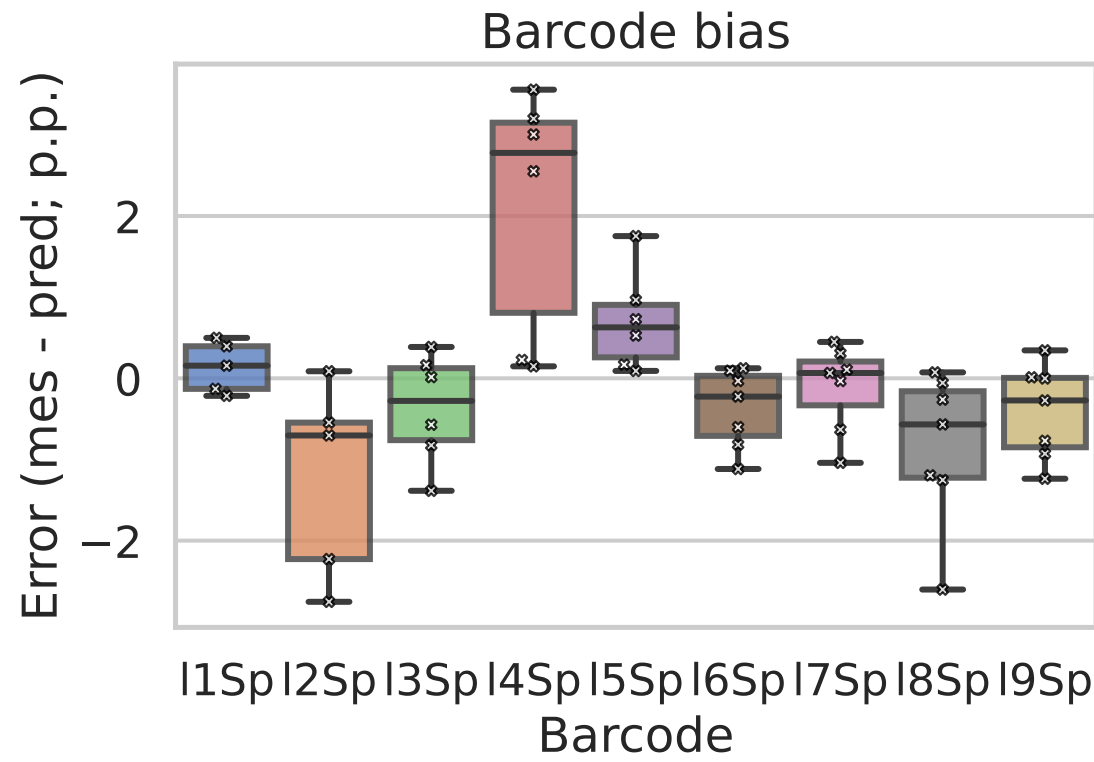
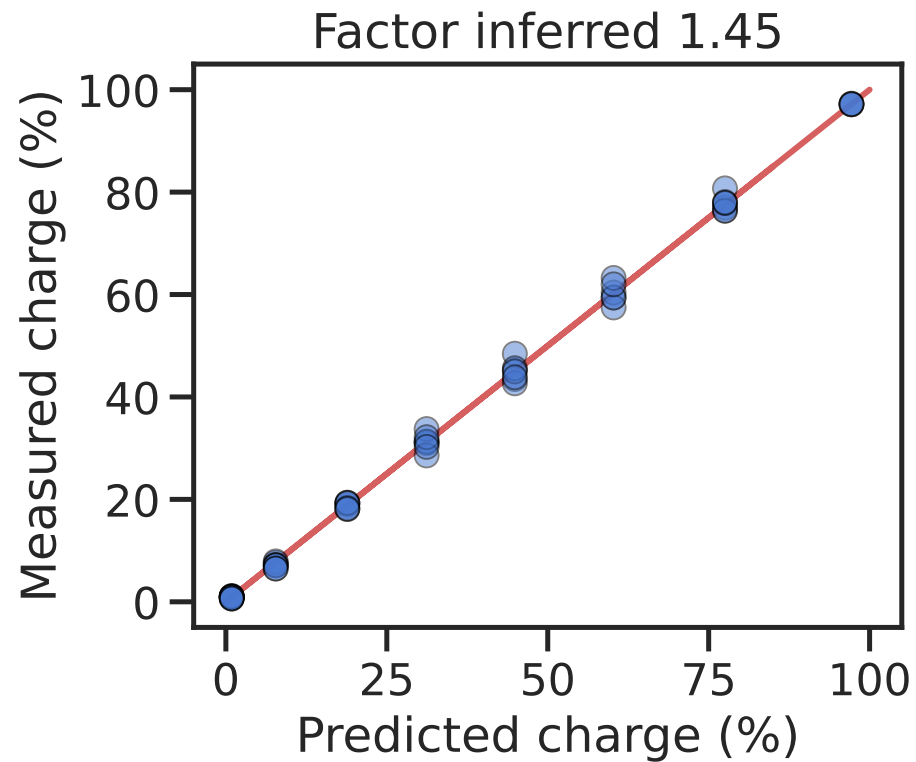
Pro-TGG-1-1



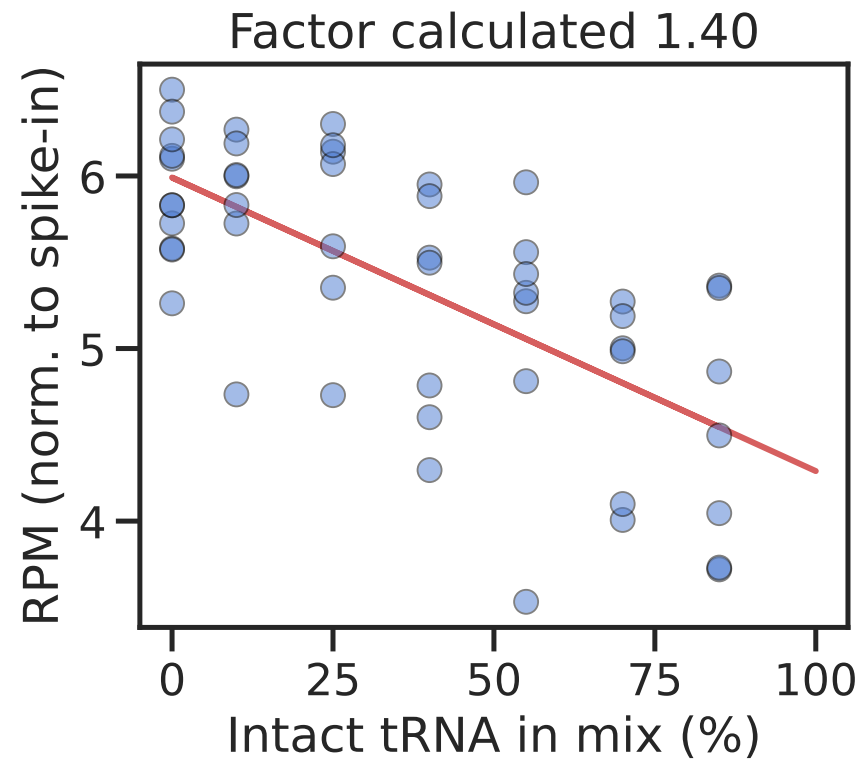
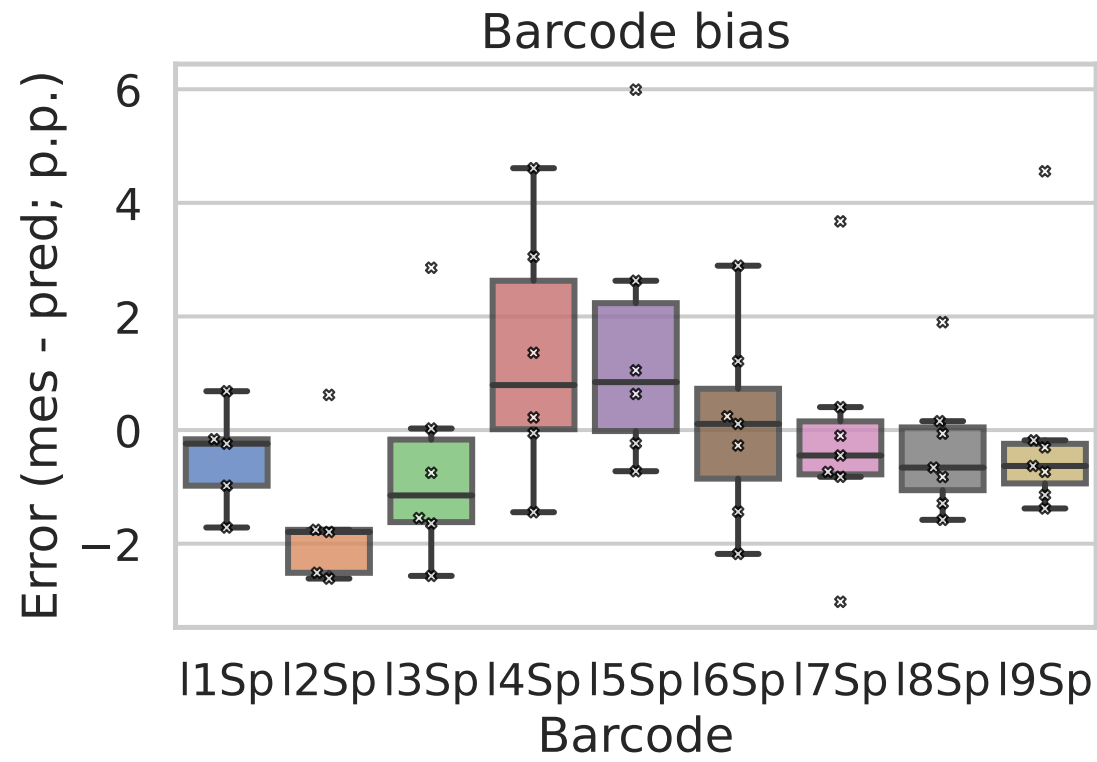
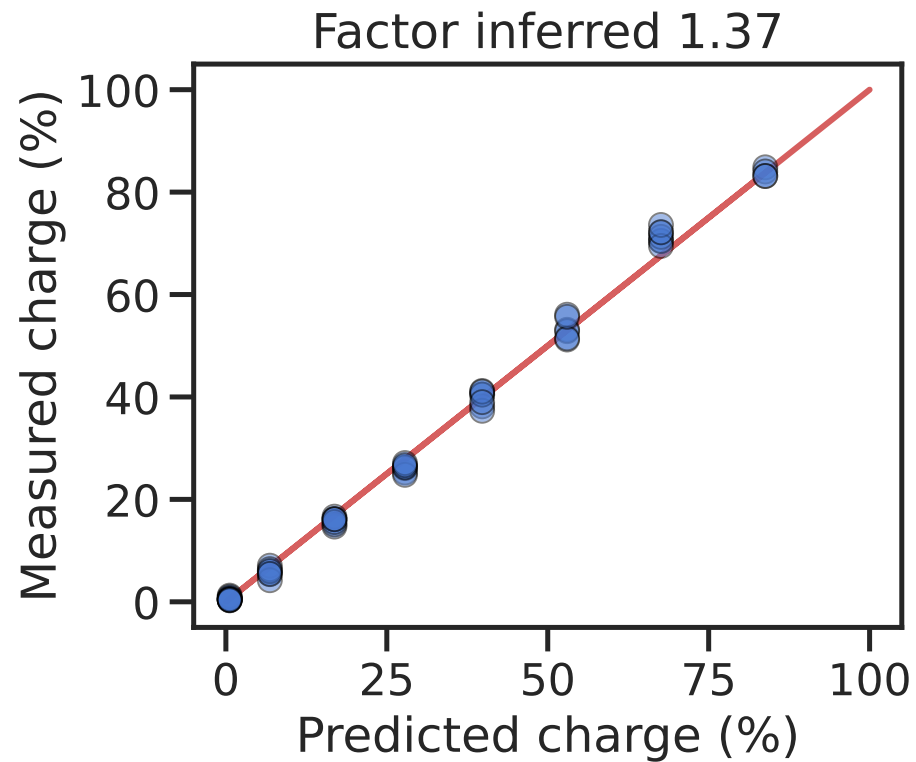
Ala-CGC-4-1



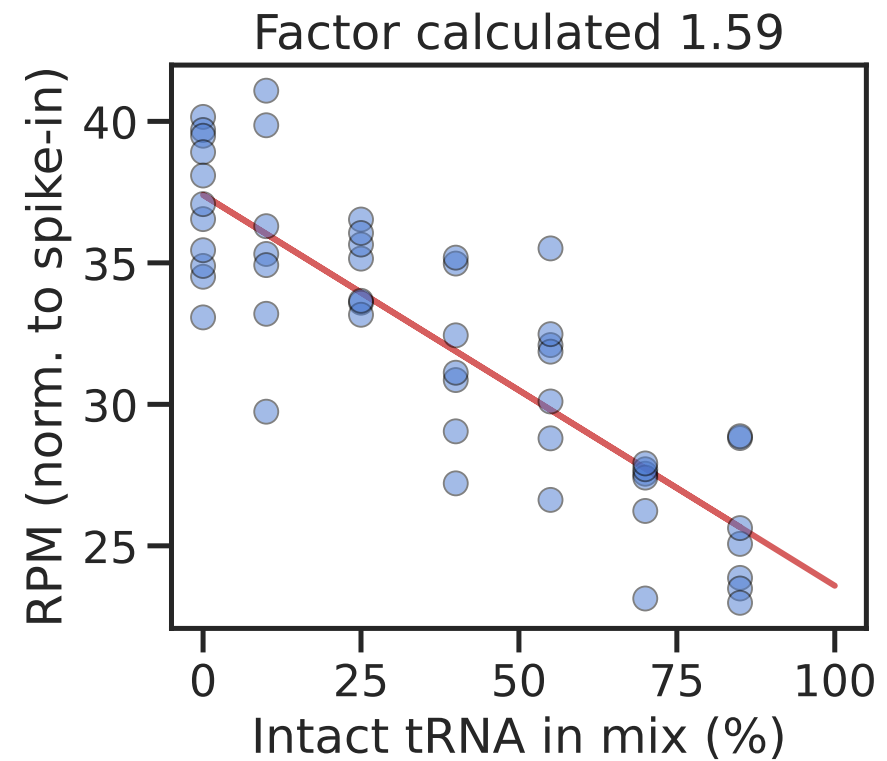
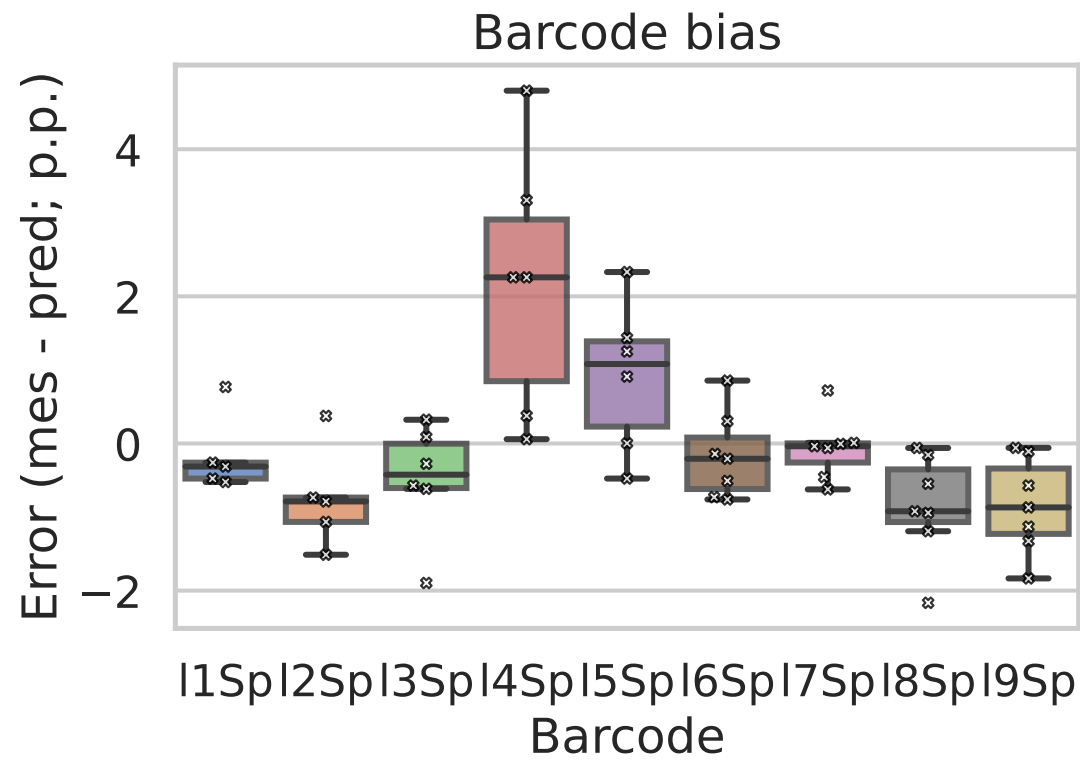
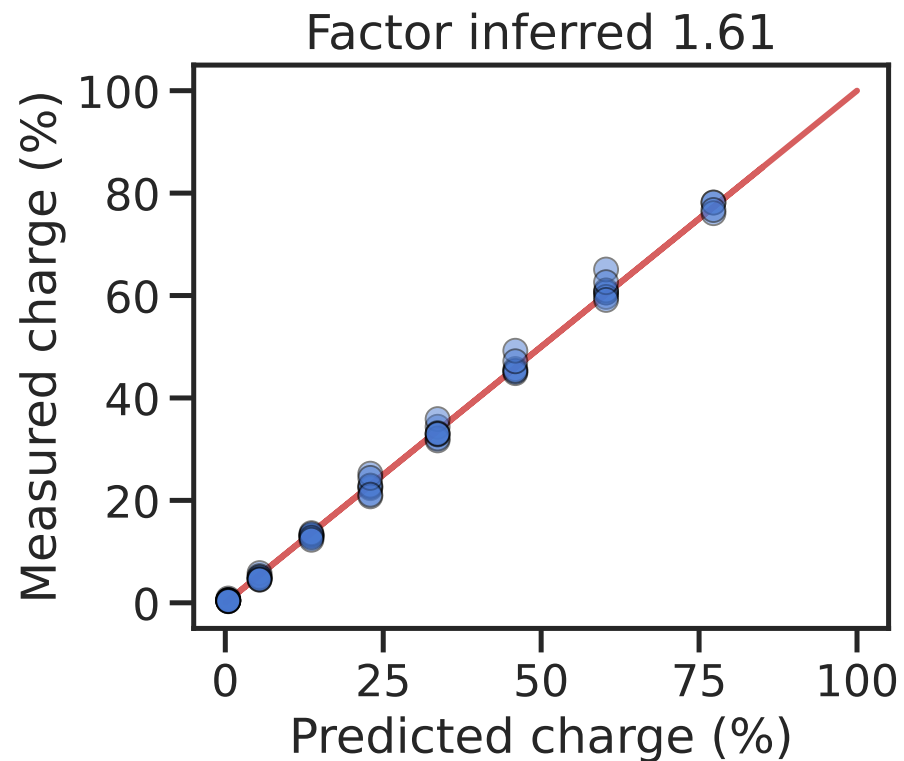
Gly-TCC-3-1



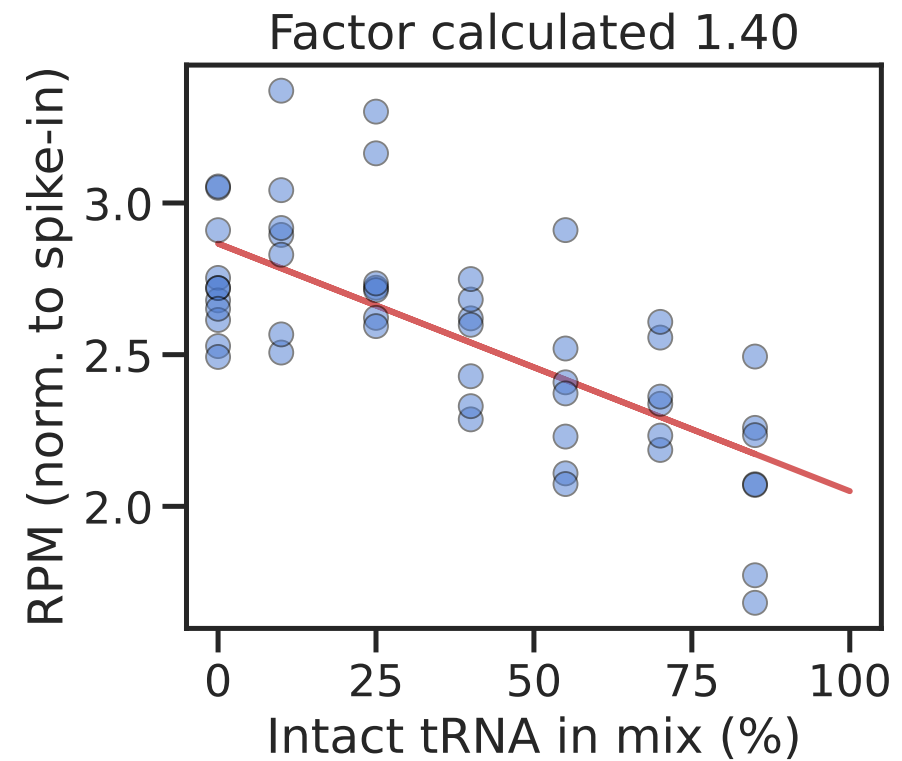
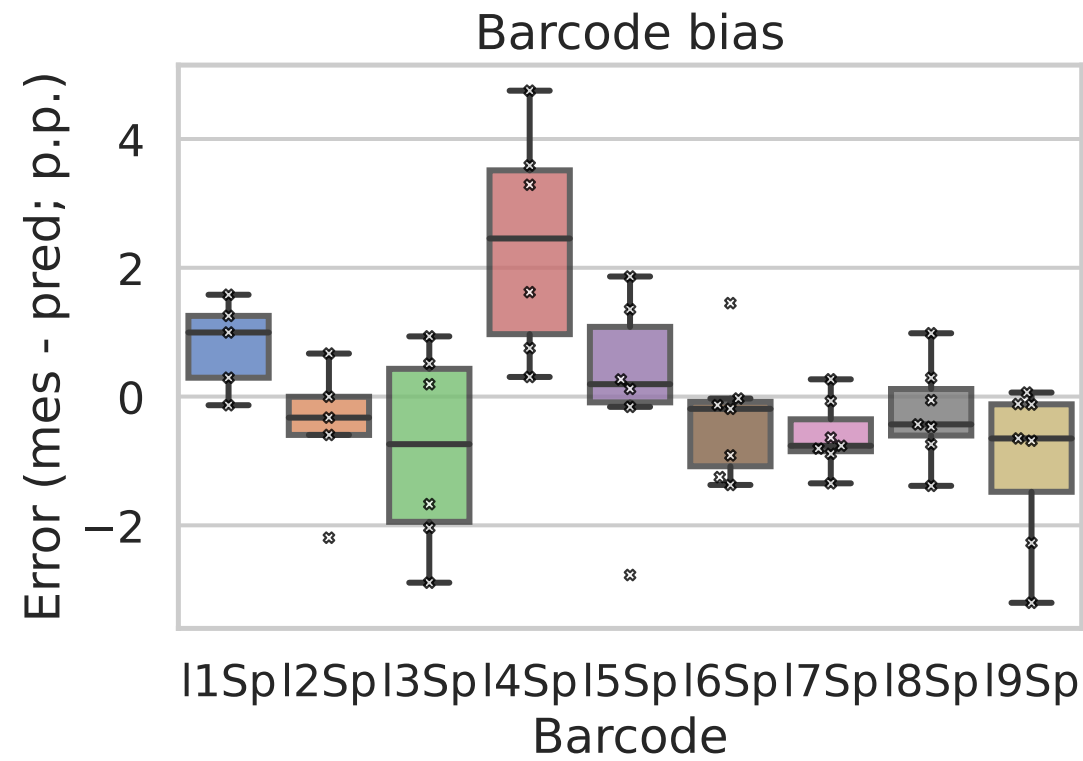
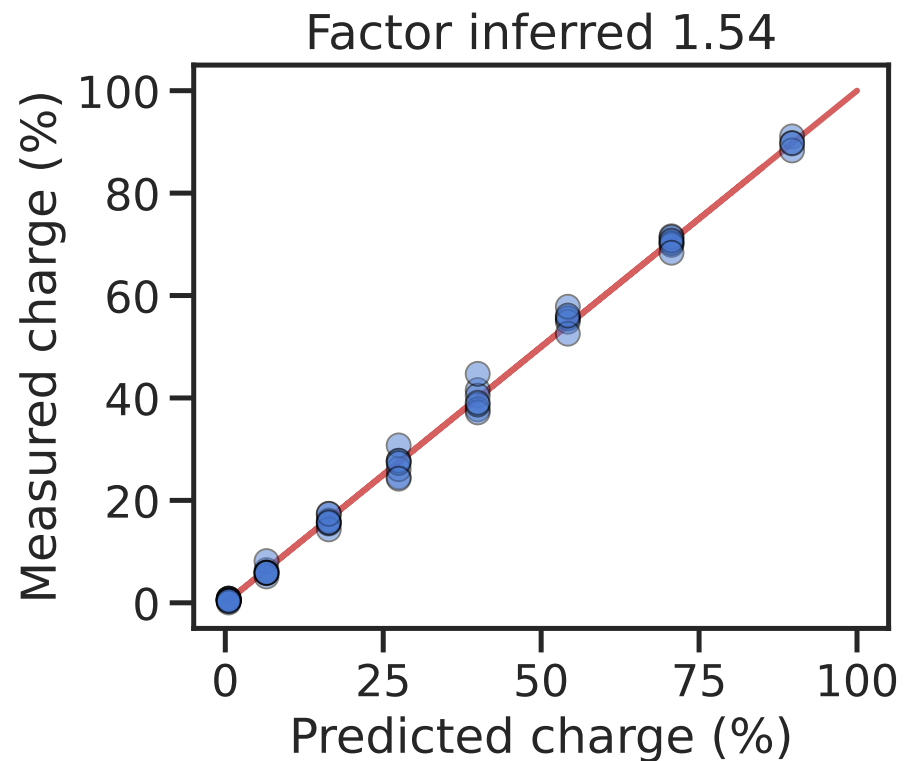
Ser-CGA-4-1



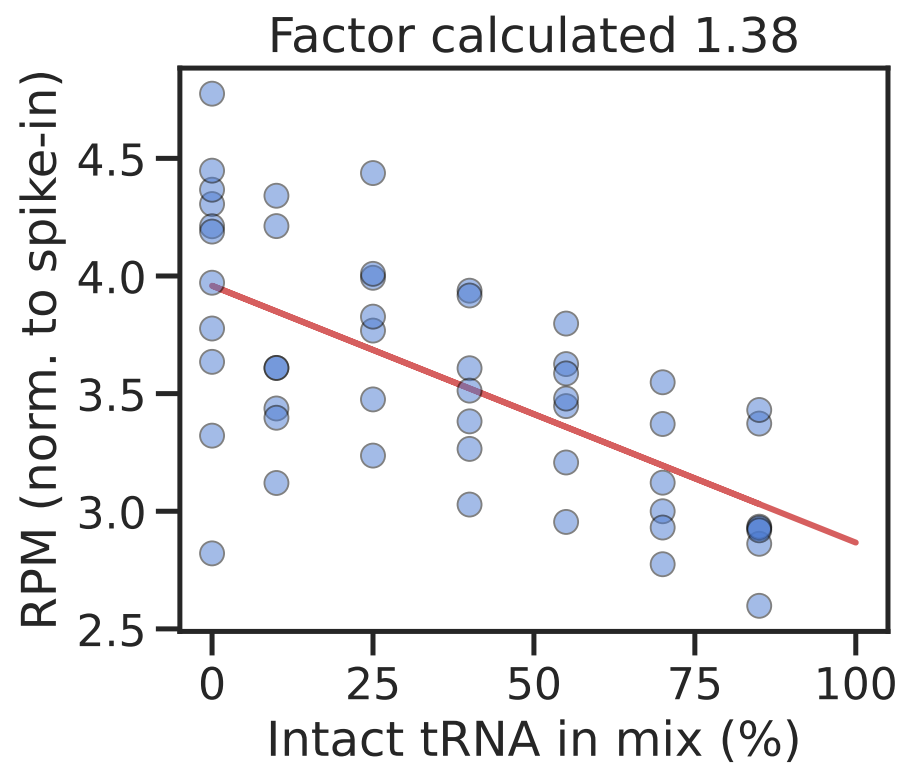
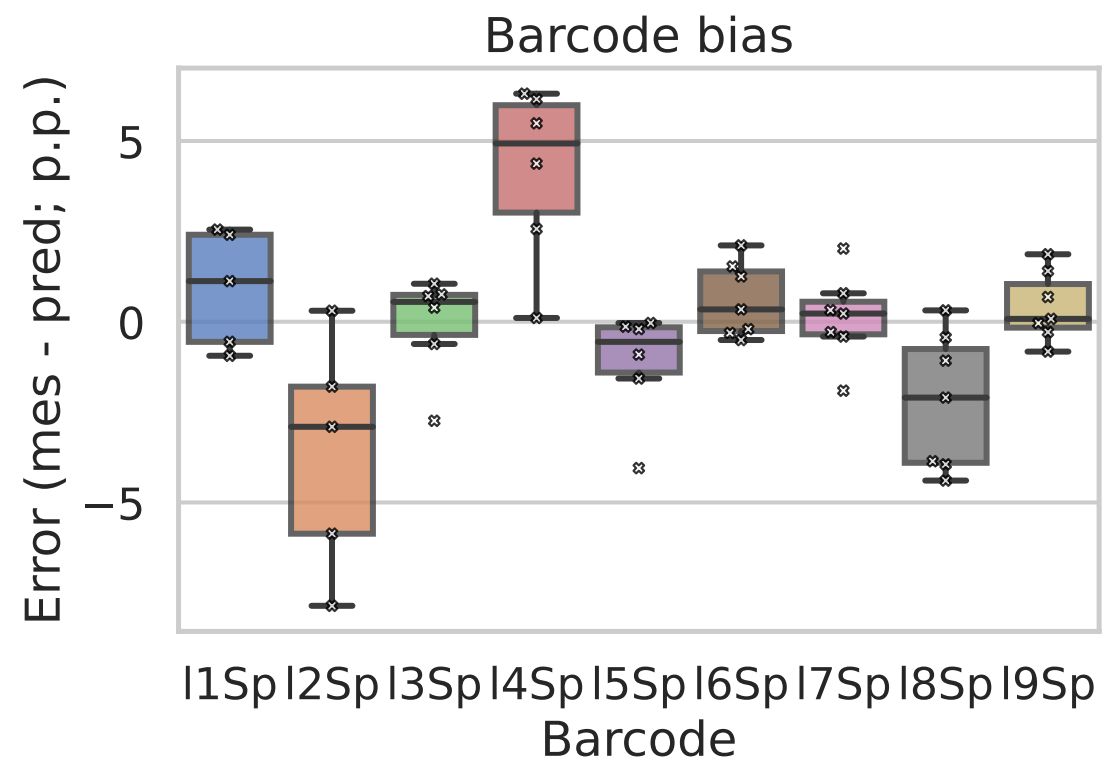
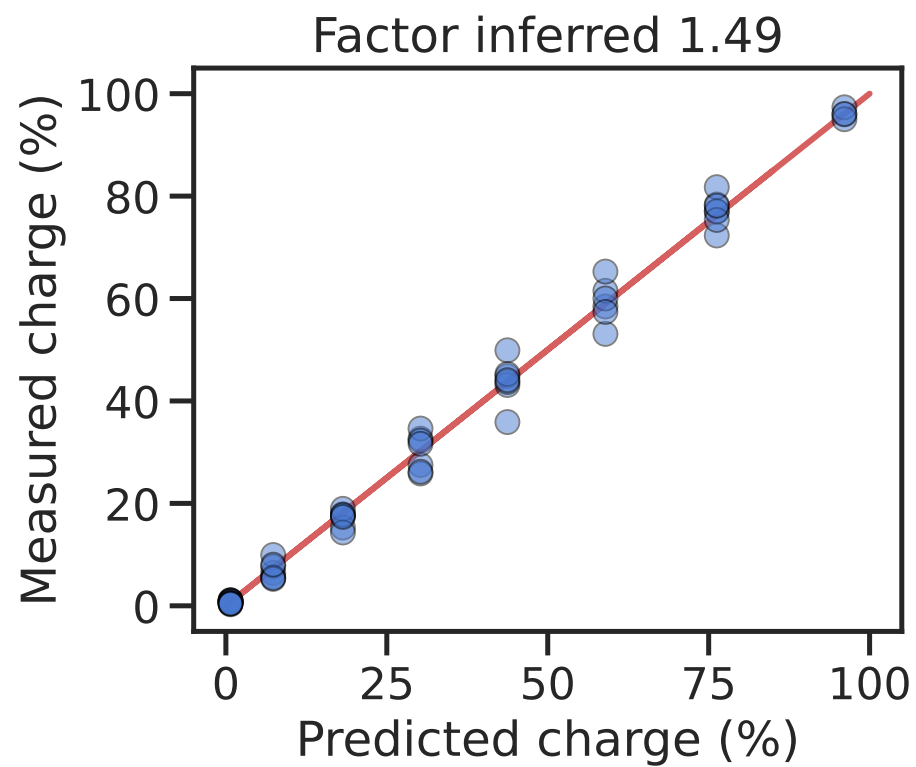
Glu-TTC-2-1



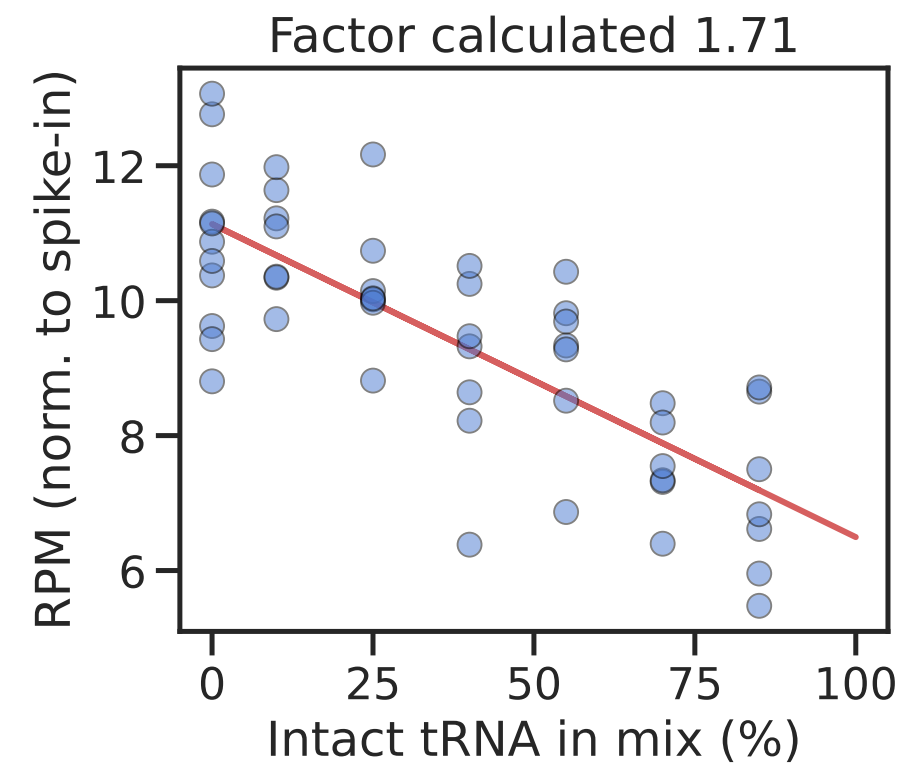
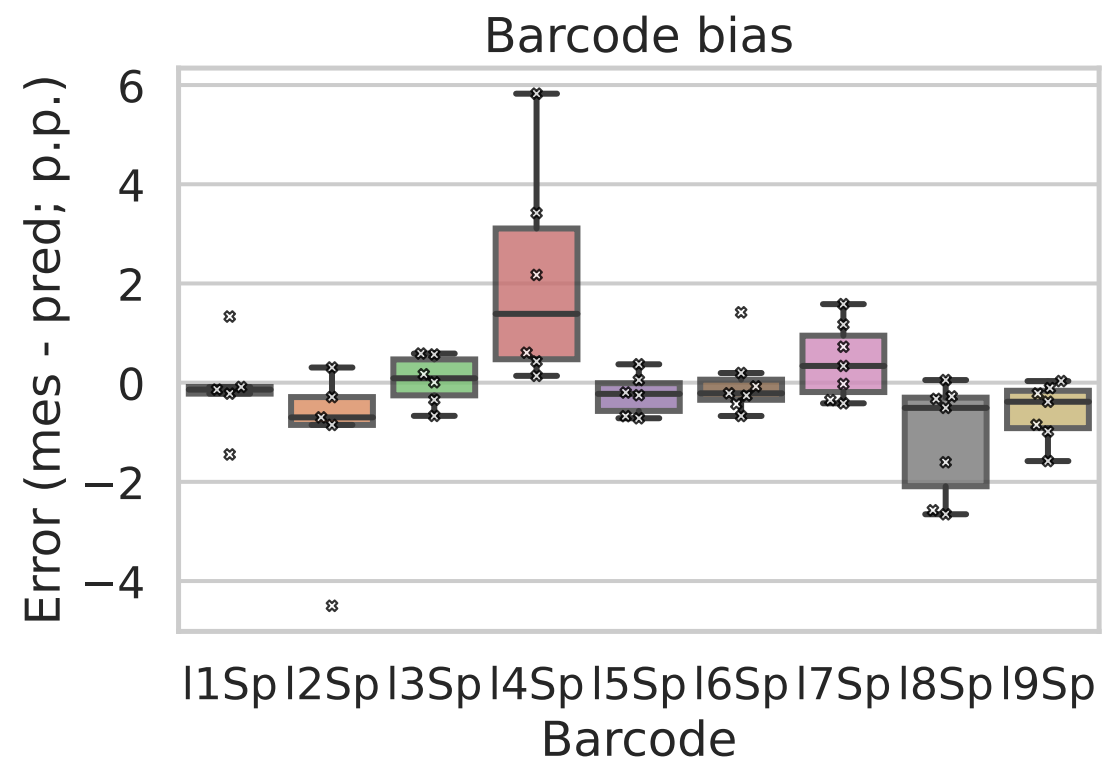
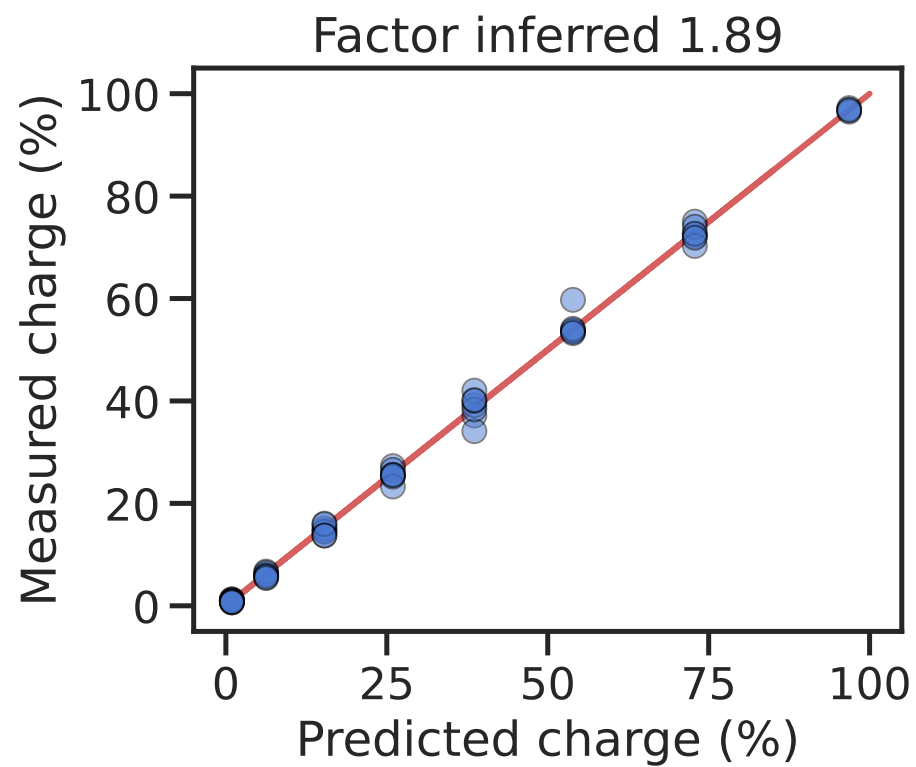
mito-Asn-GTT



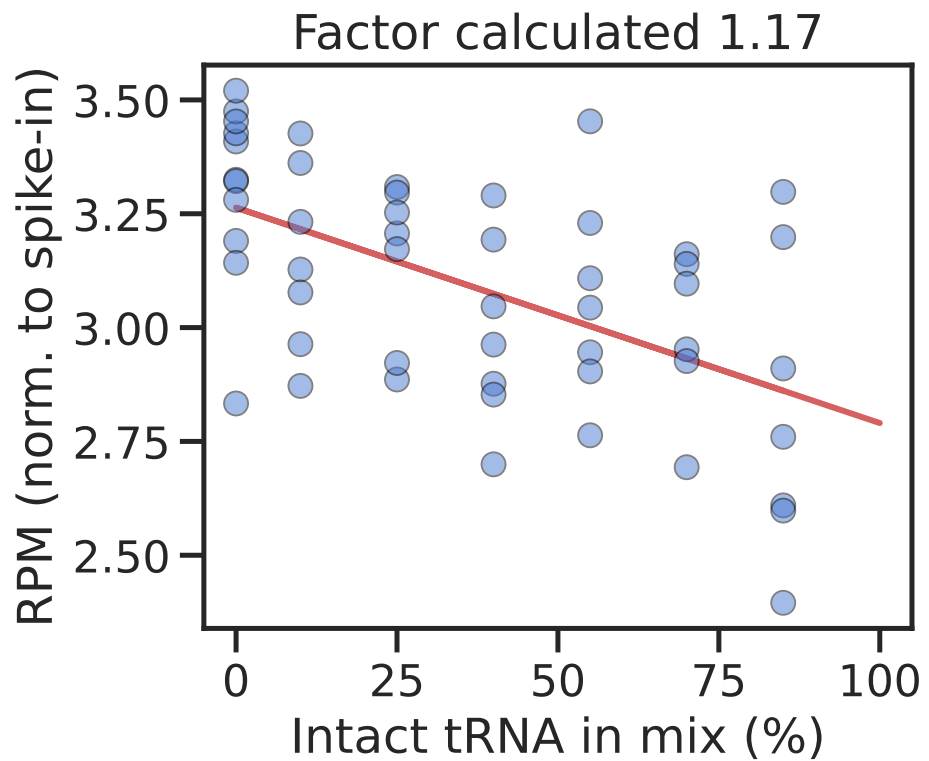
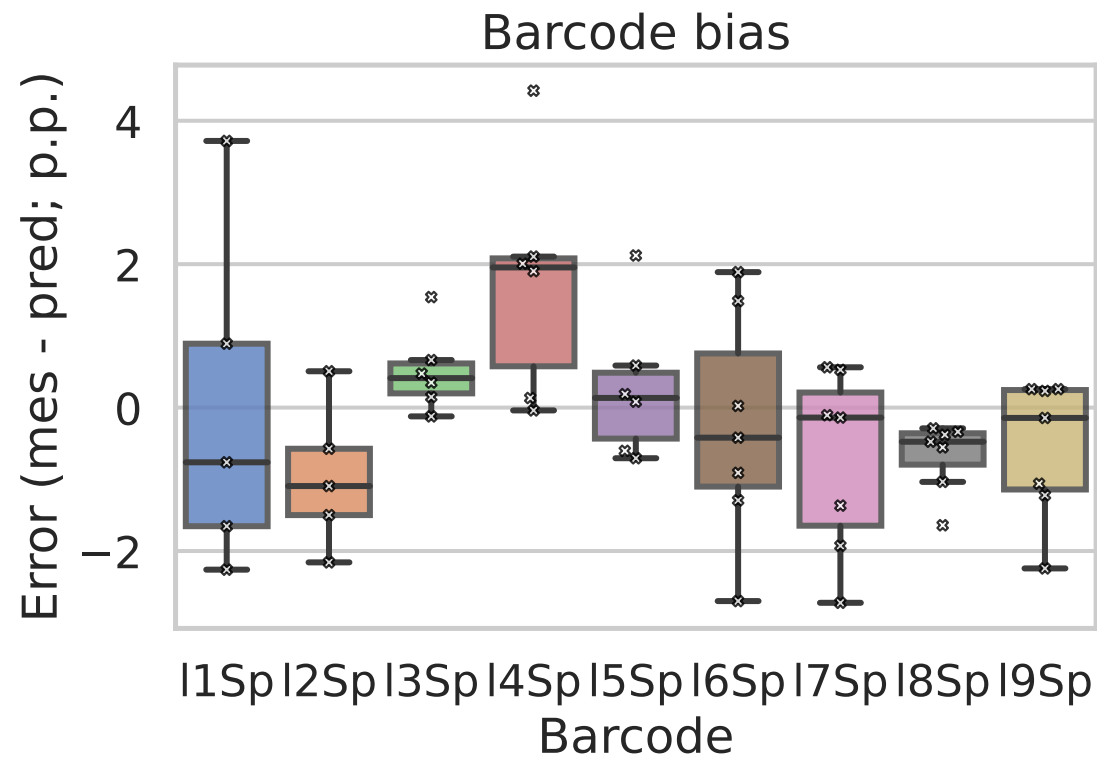
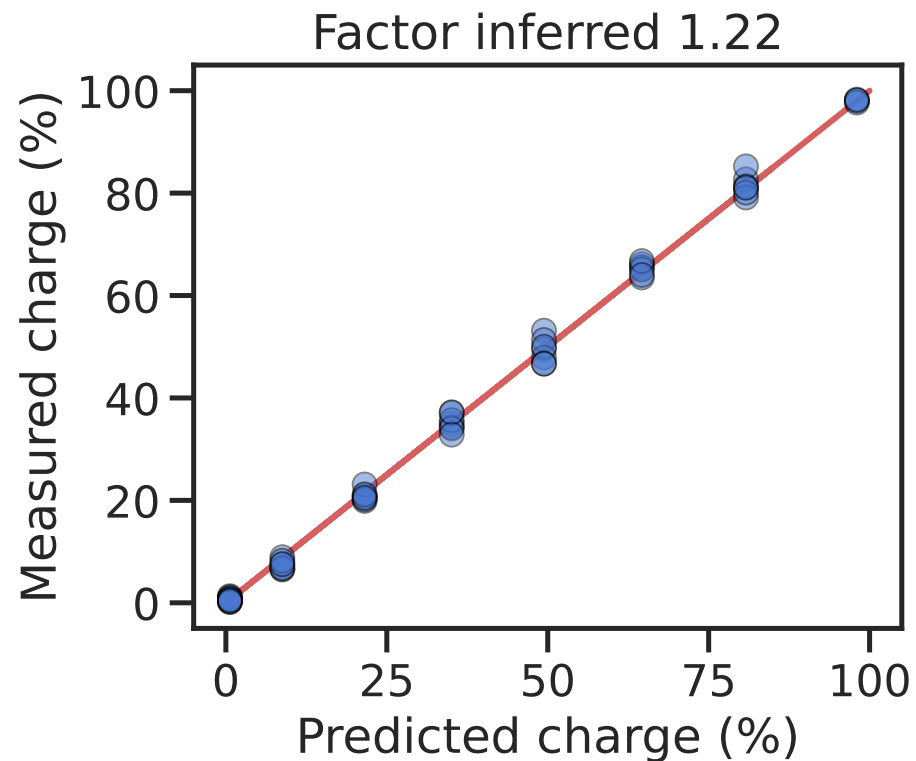
Thr-CGT-4-1



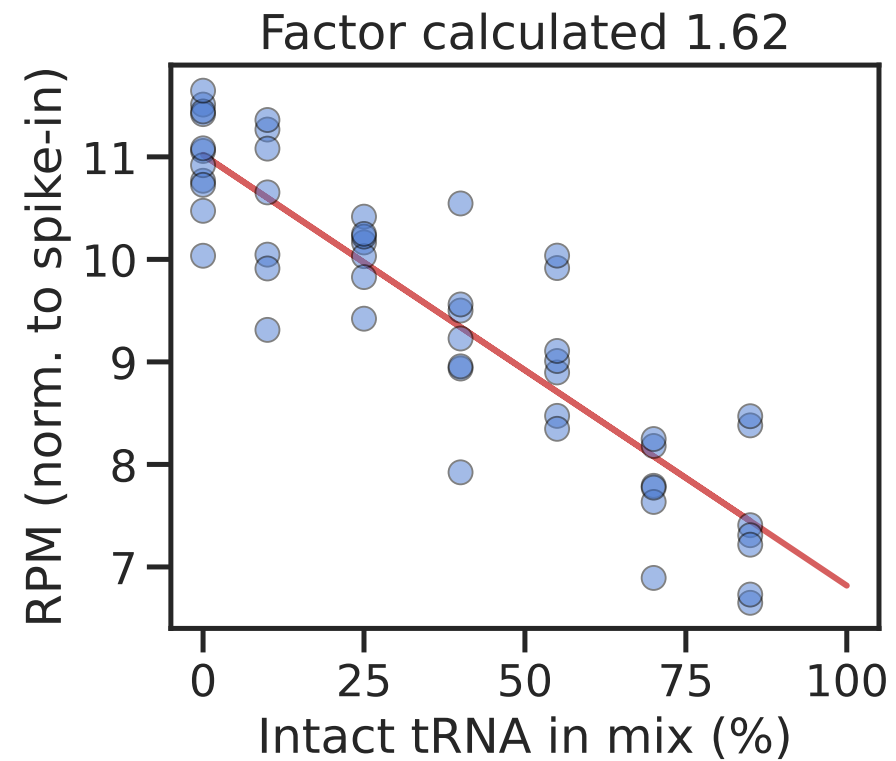
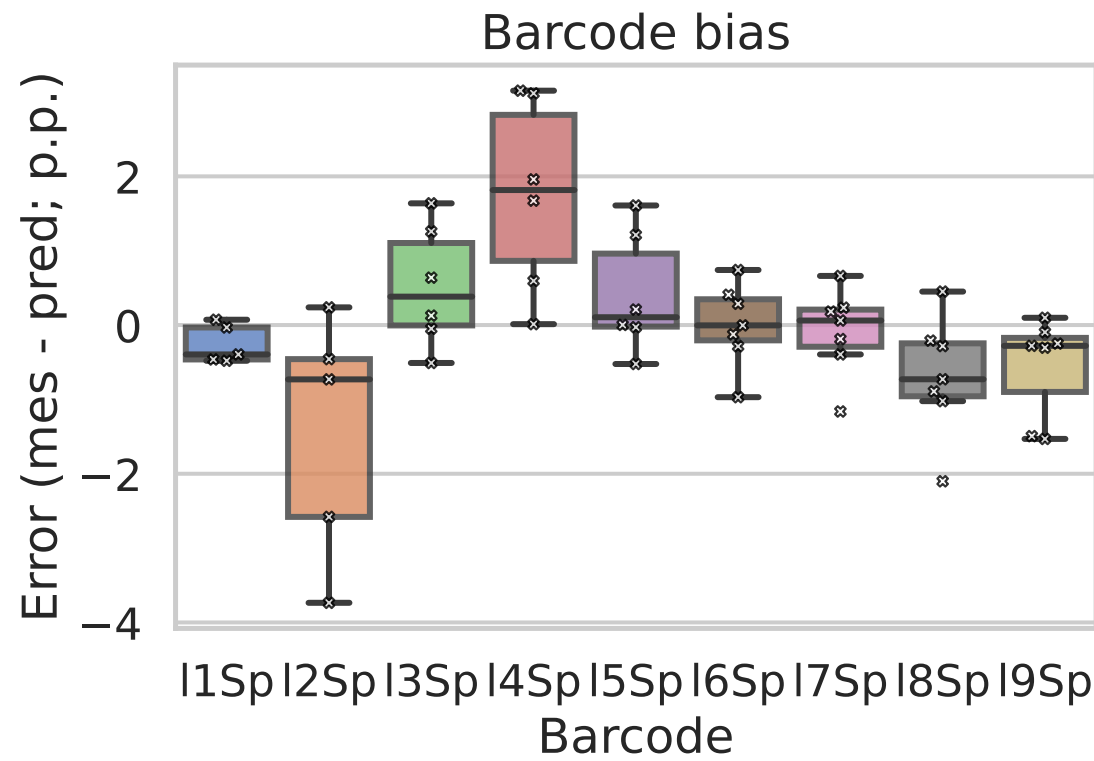
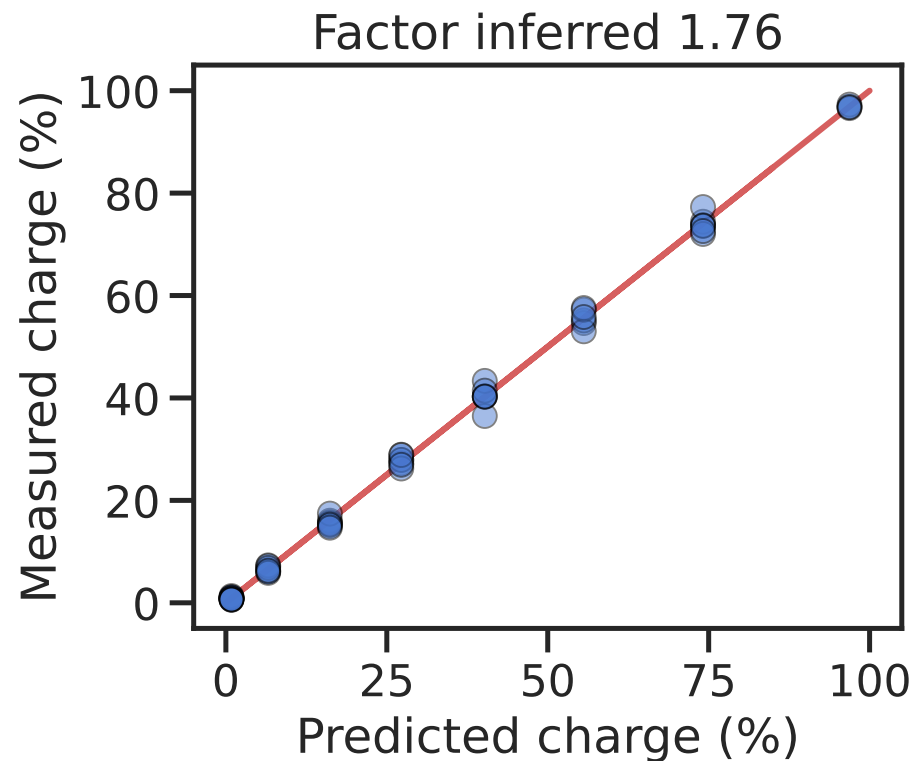
Val-AAC-2-1



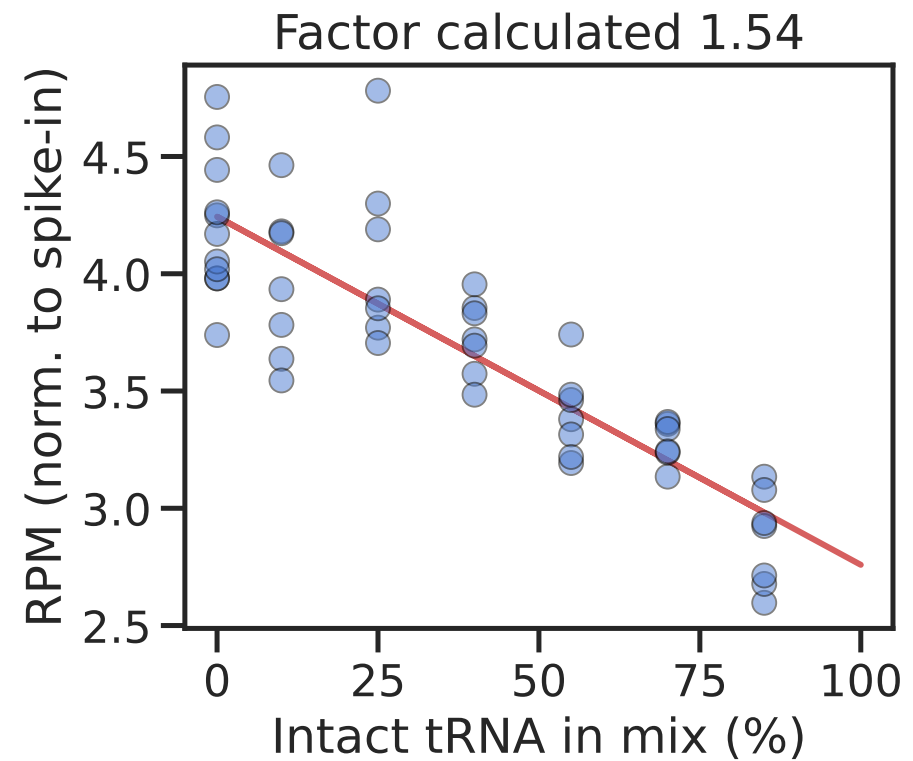
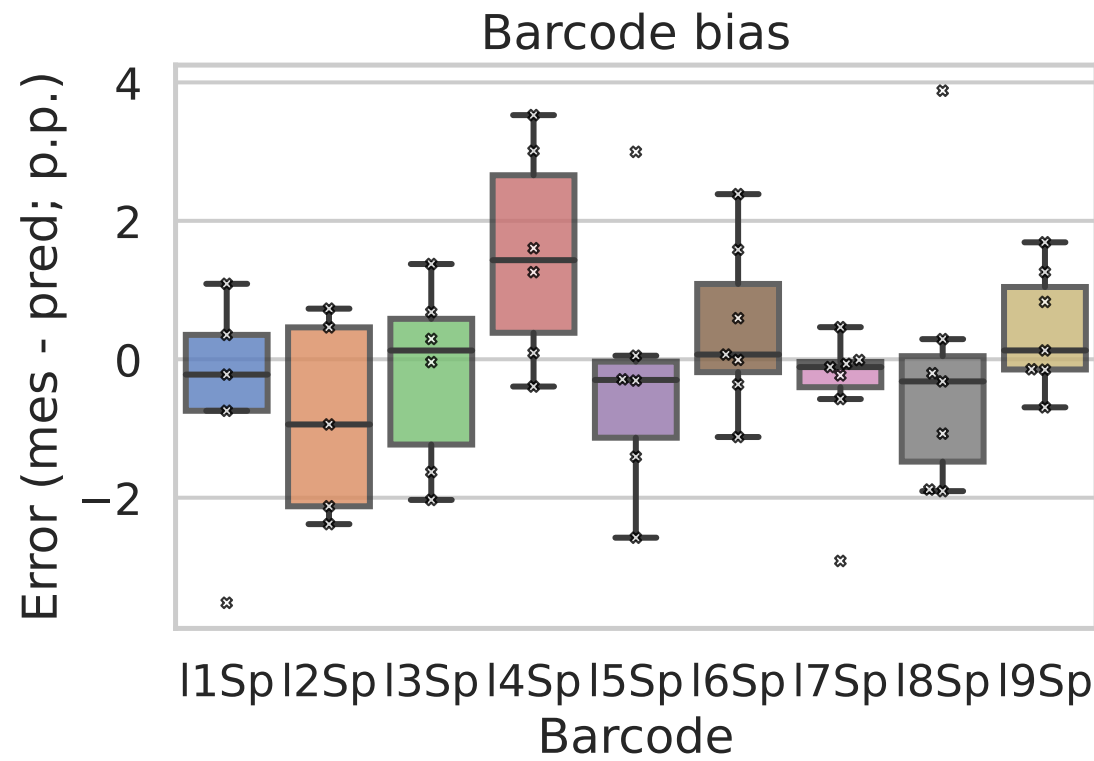
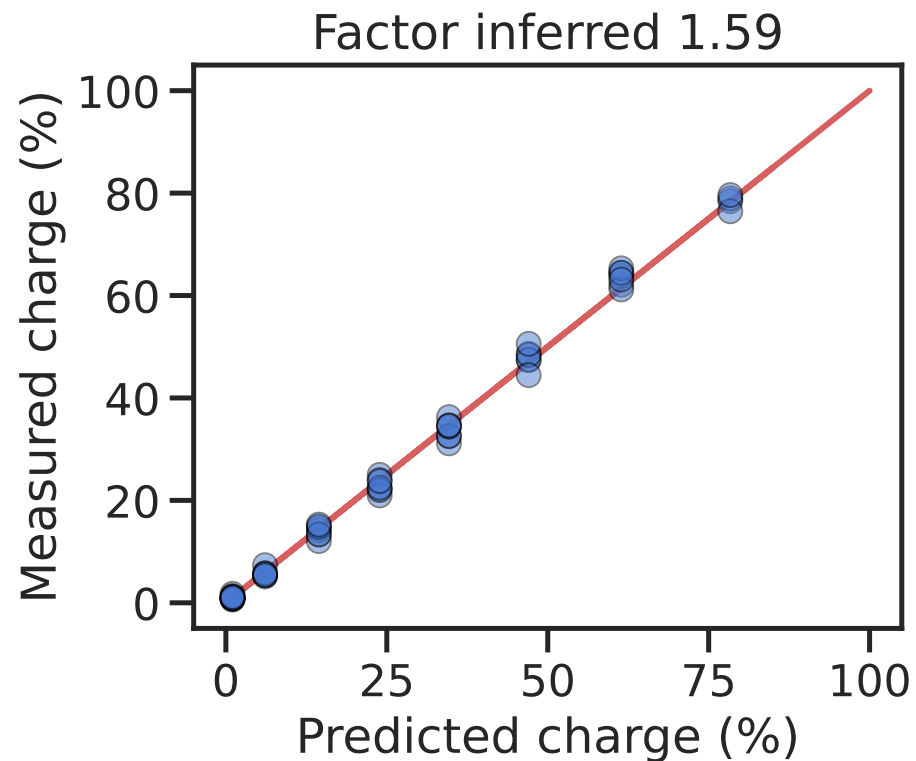
Ala-TGC-6-1



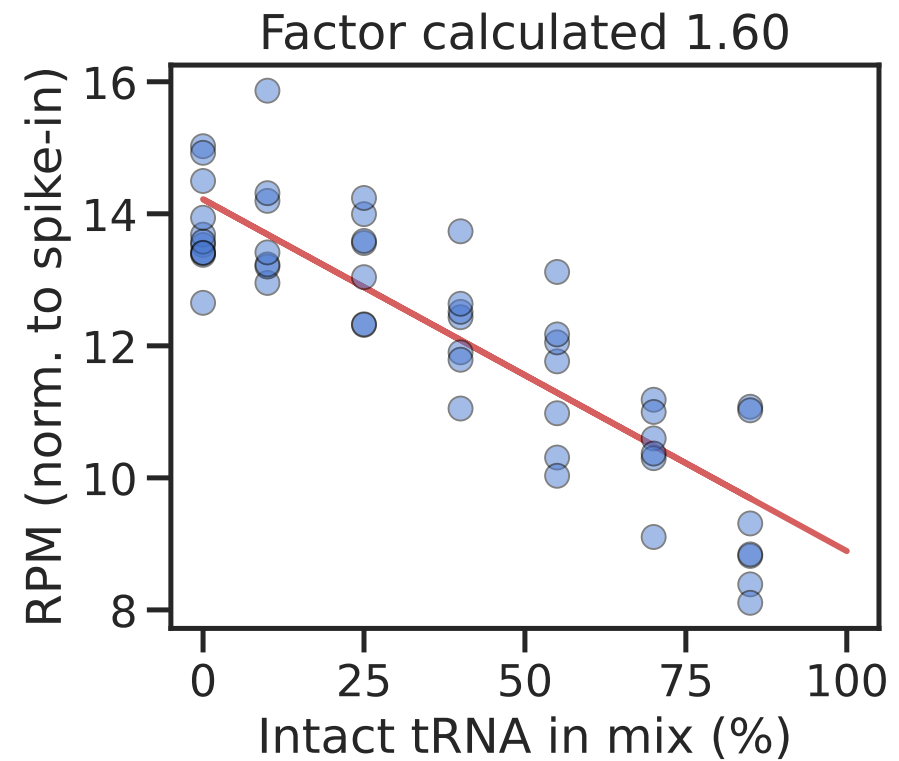
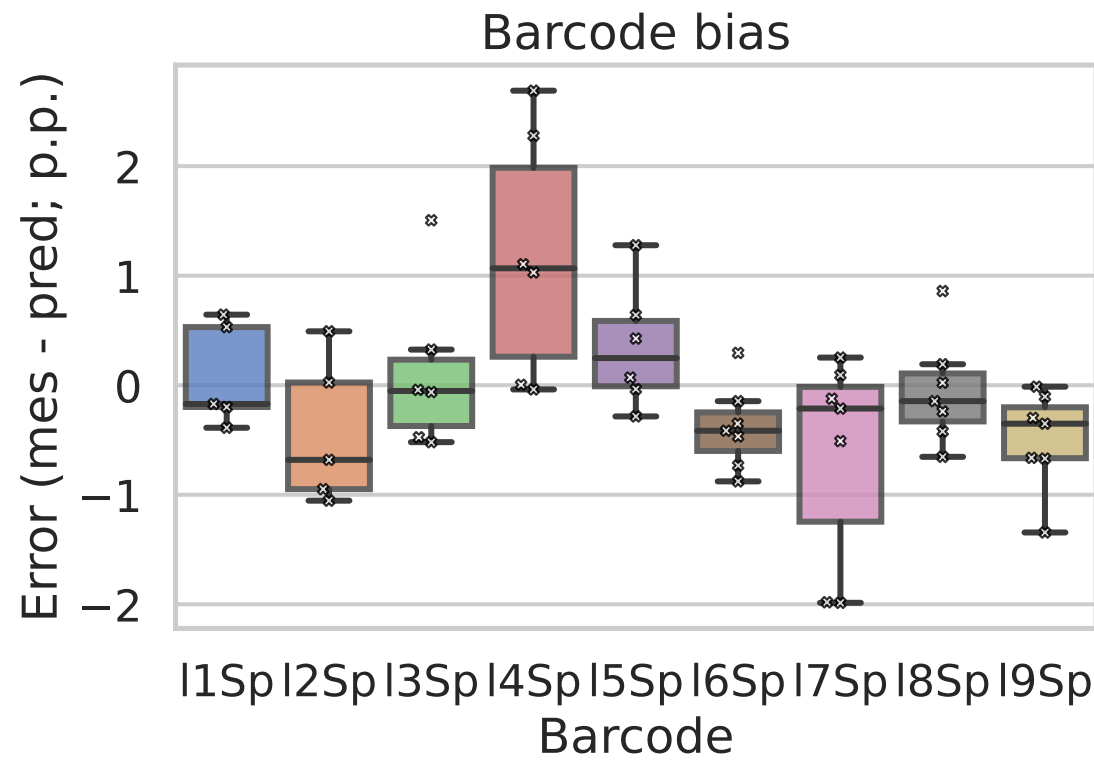
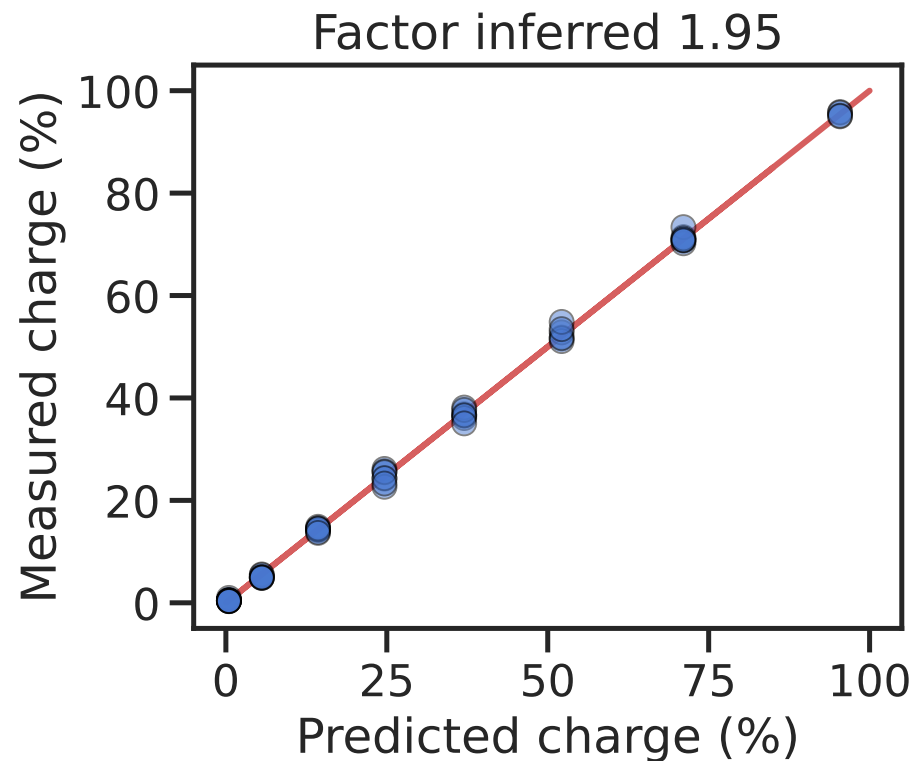
Val-CAC-3-1



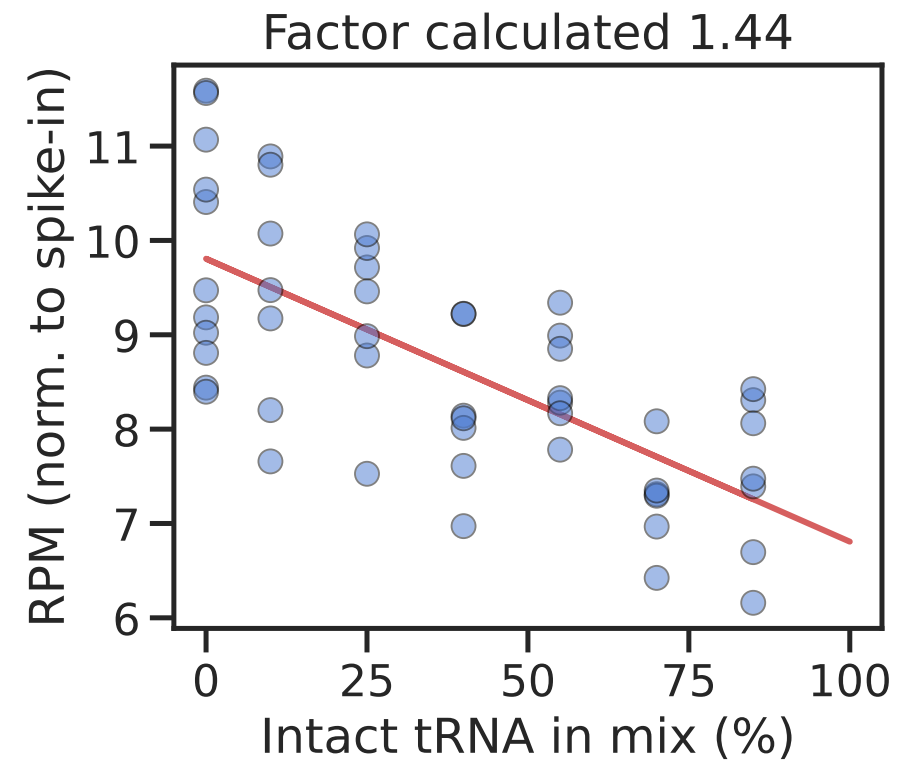
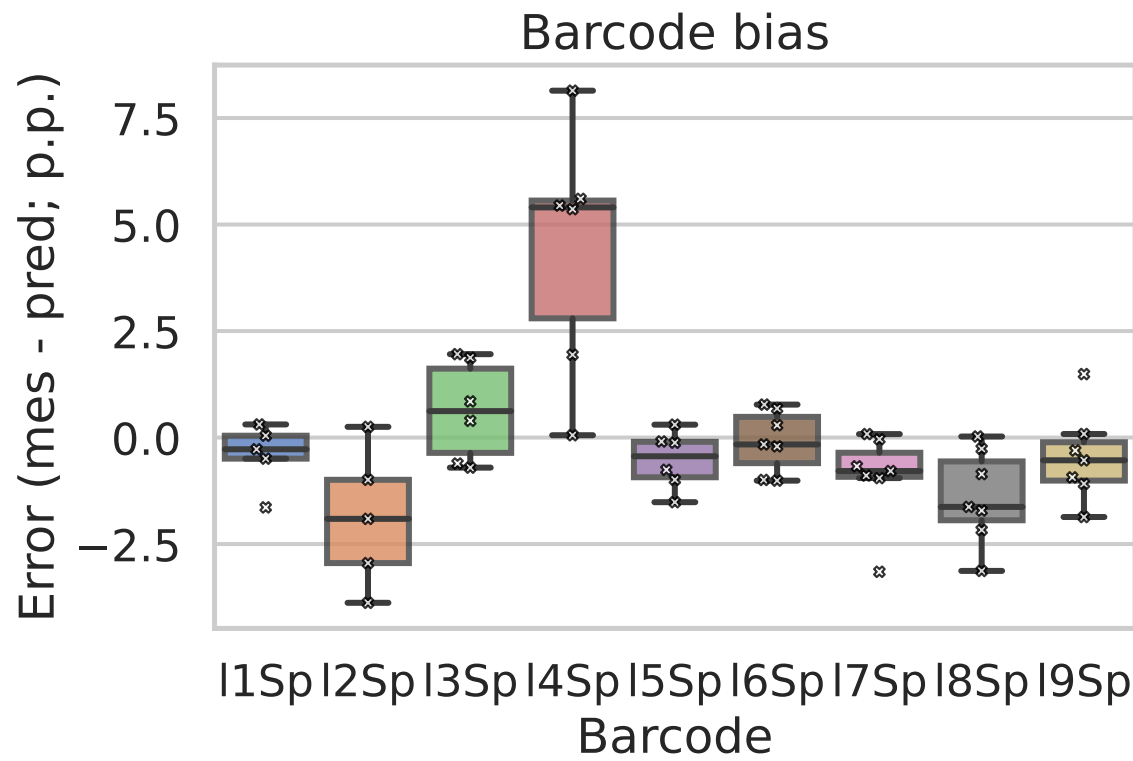
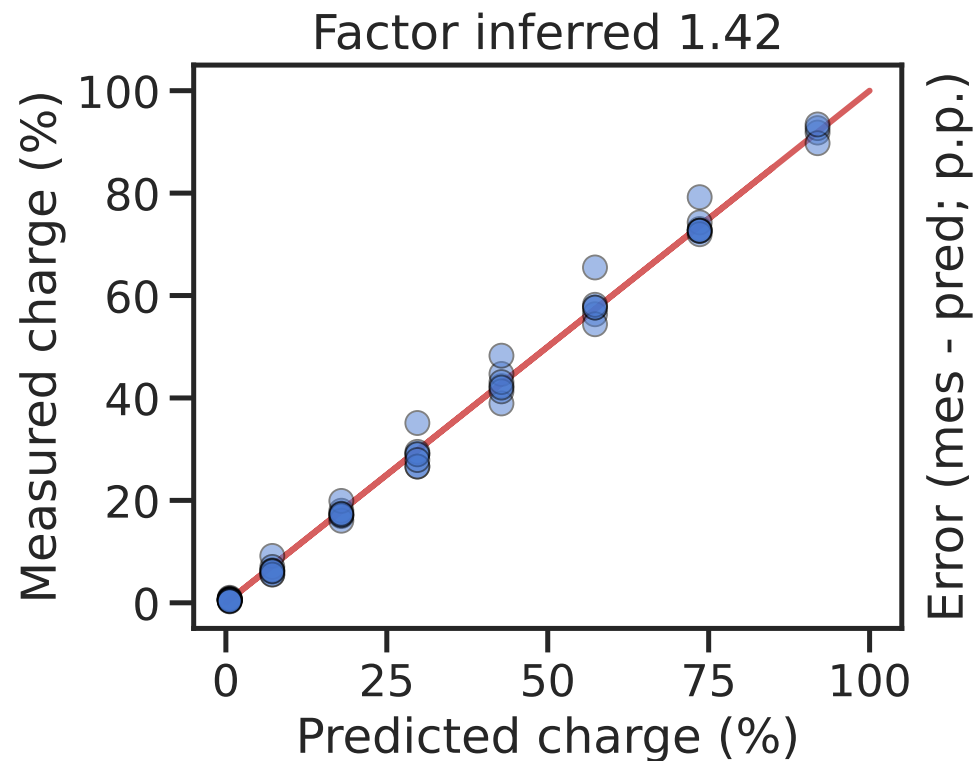
mito-Thr-TGT



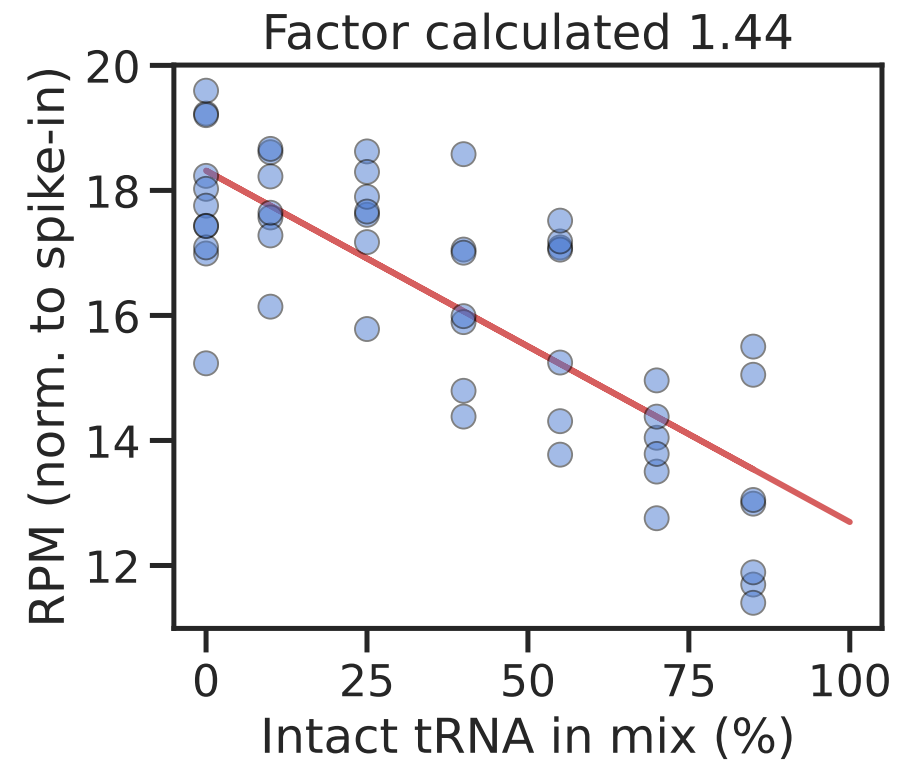
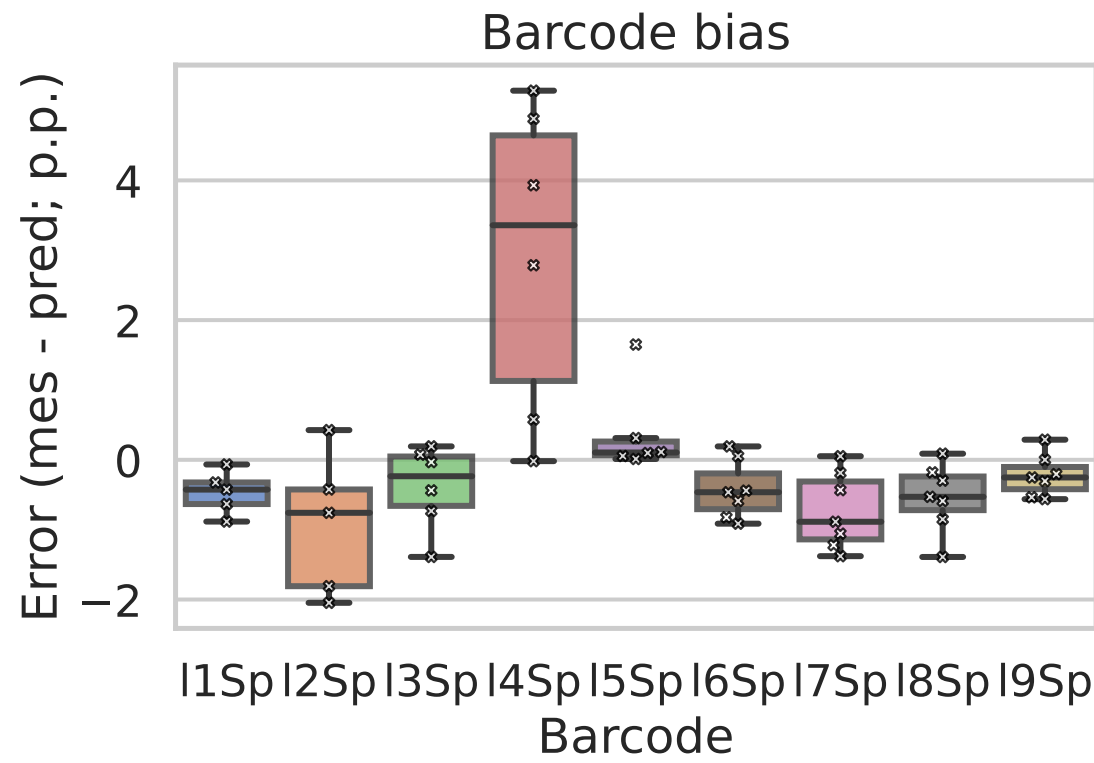
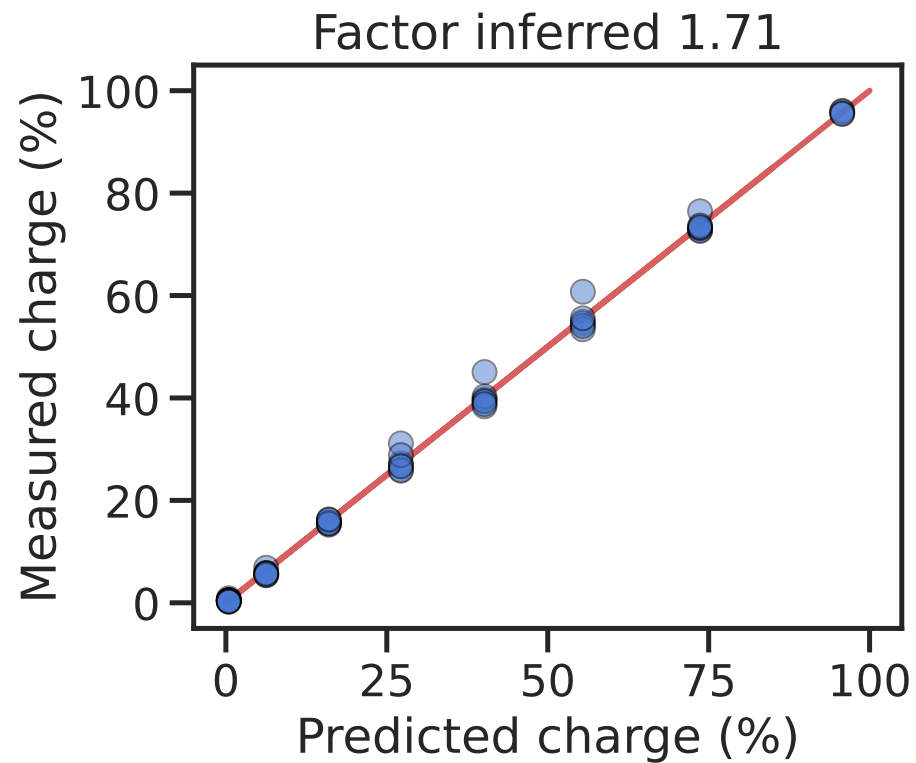
Arg-TCG-5-1



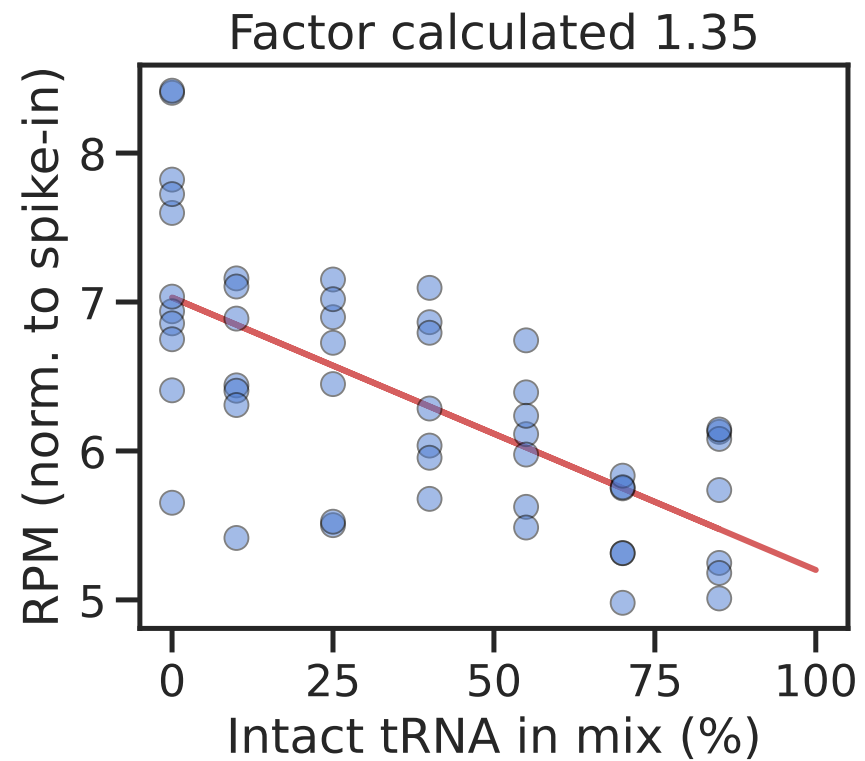
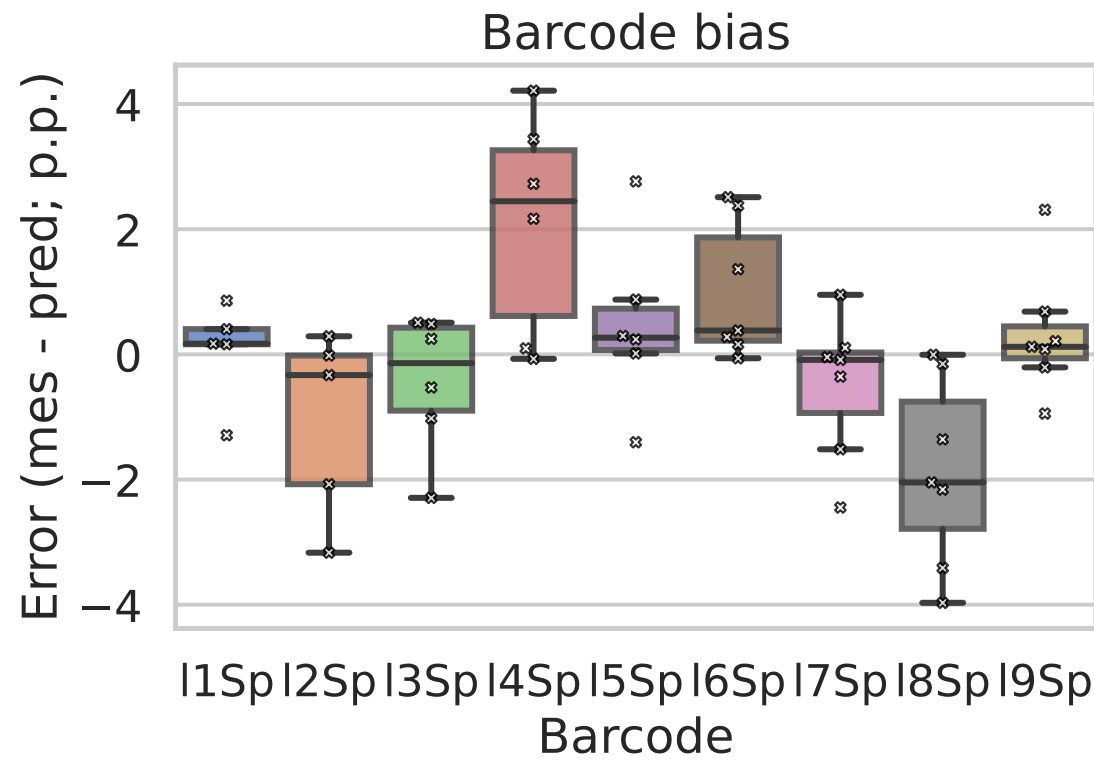
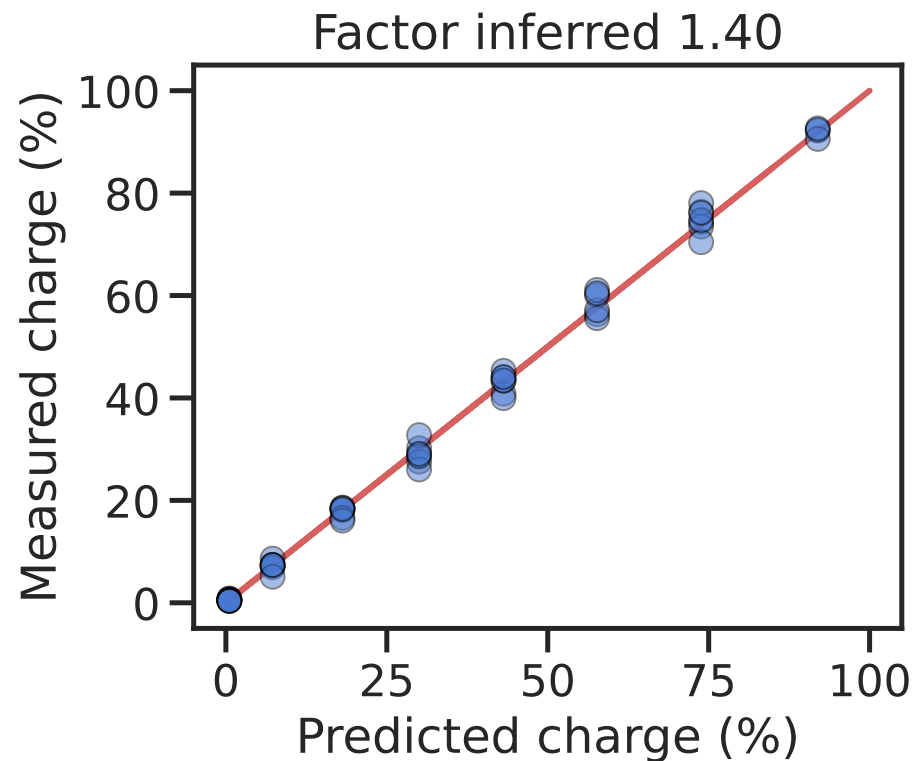
Thr-AGT-2-1



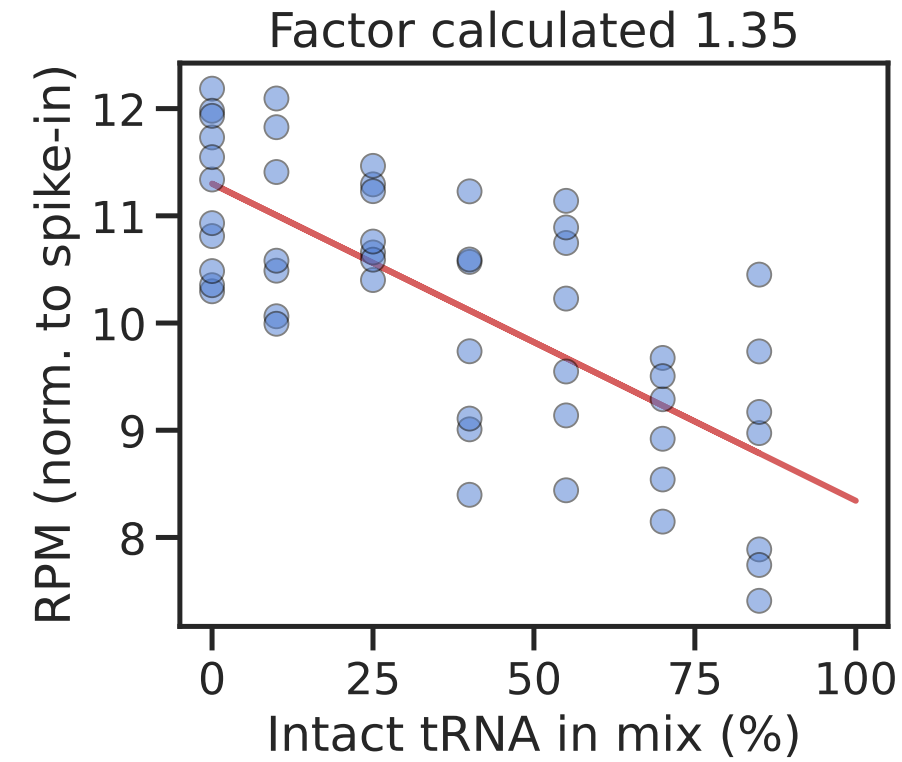
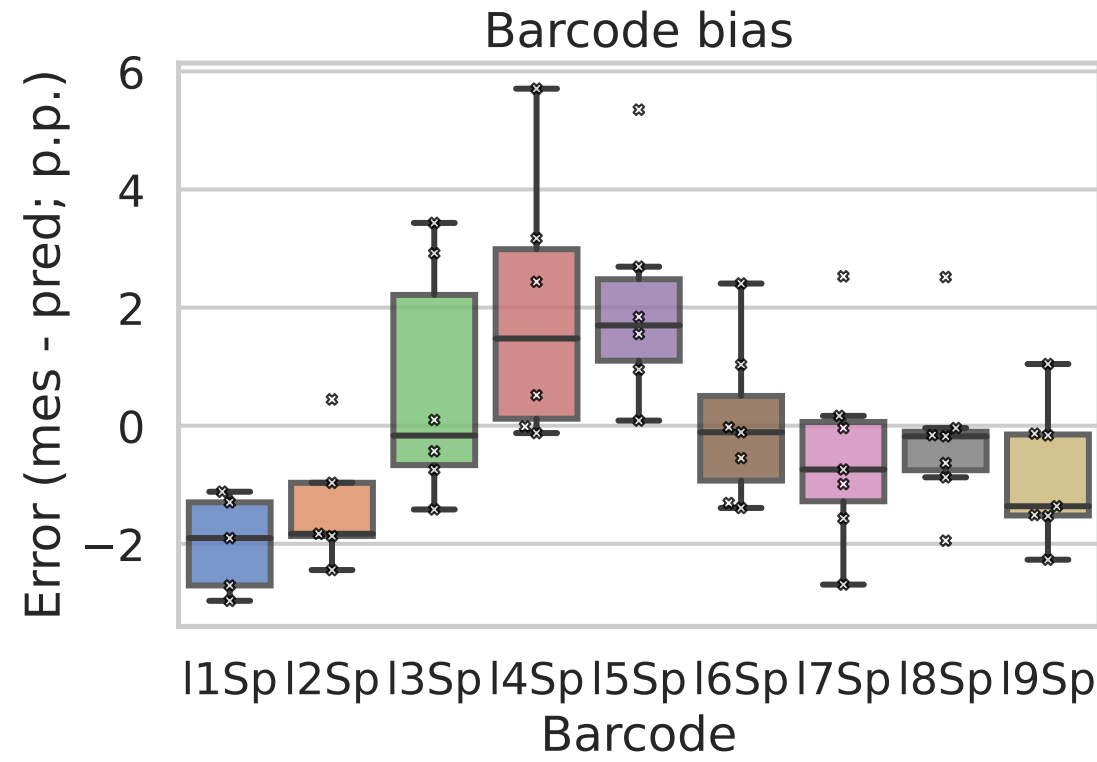
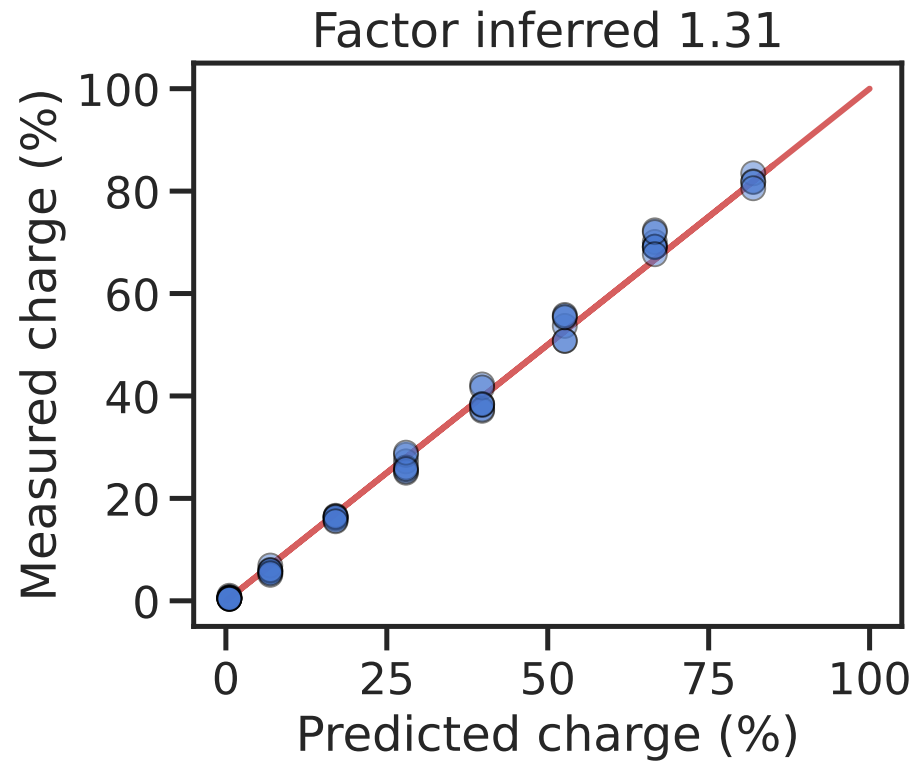
Arg-CCT-1-1



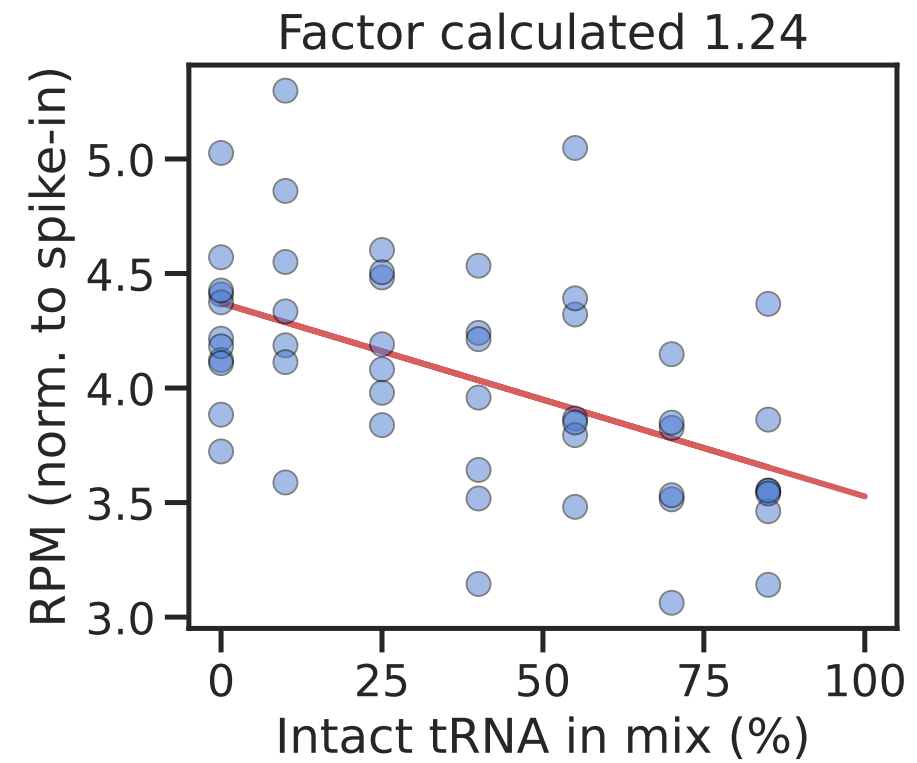
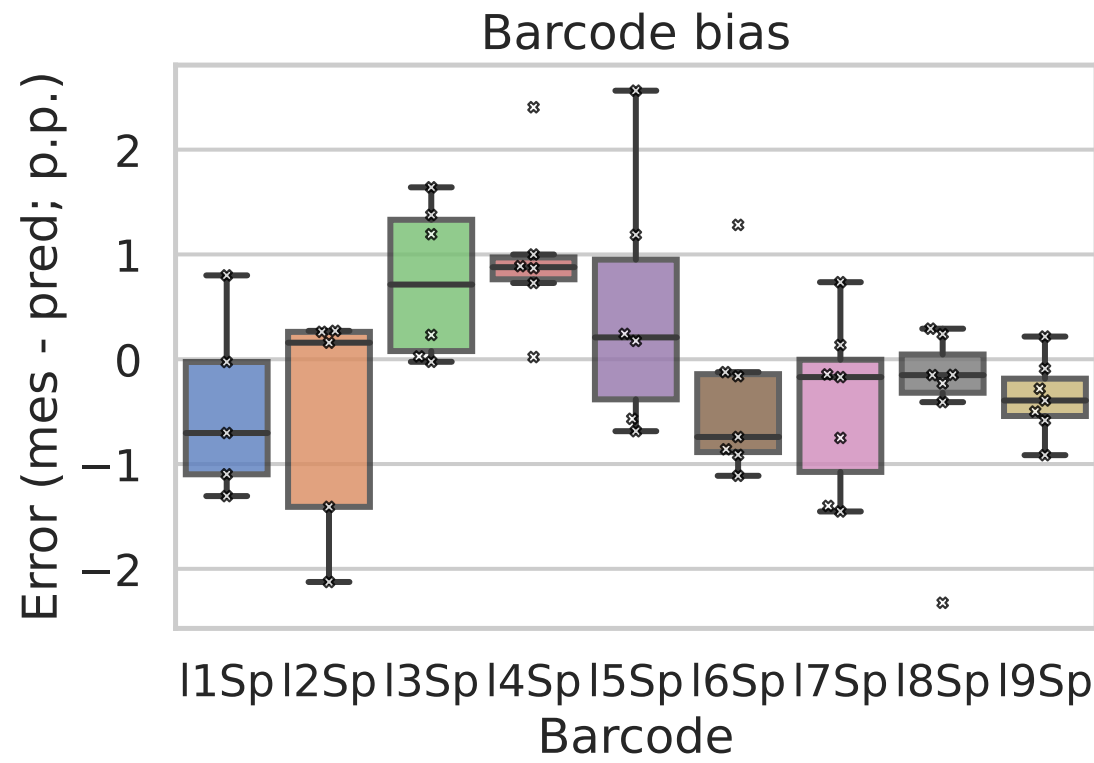
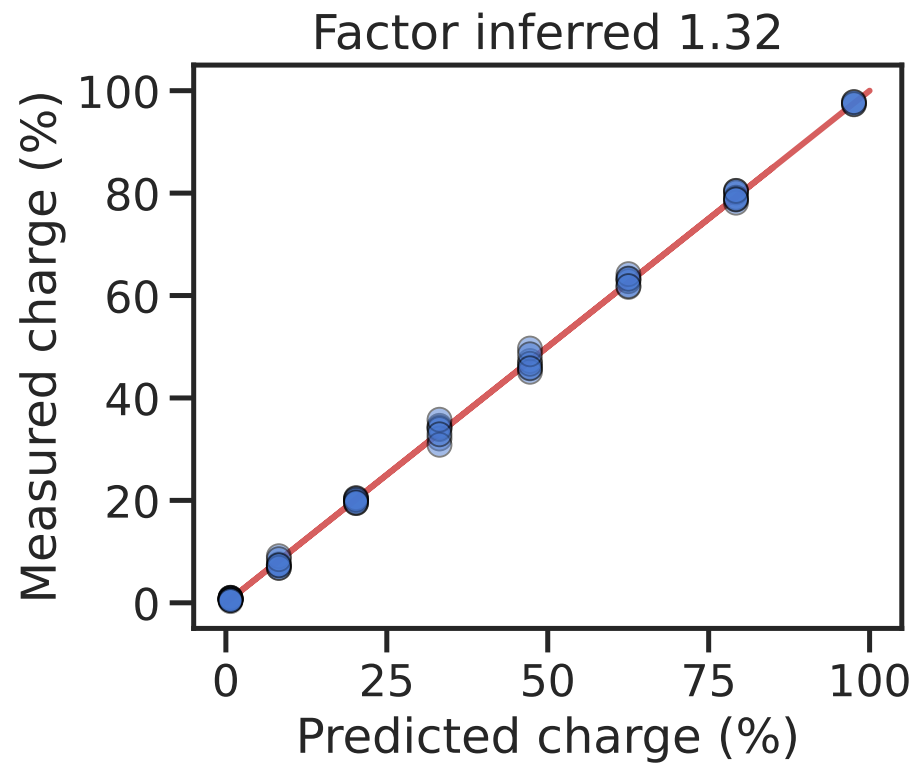
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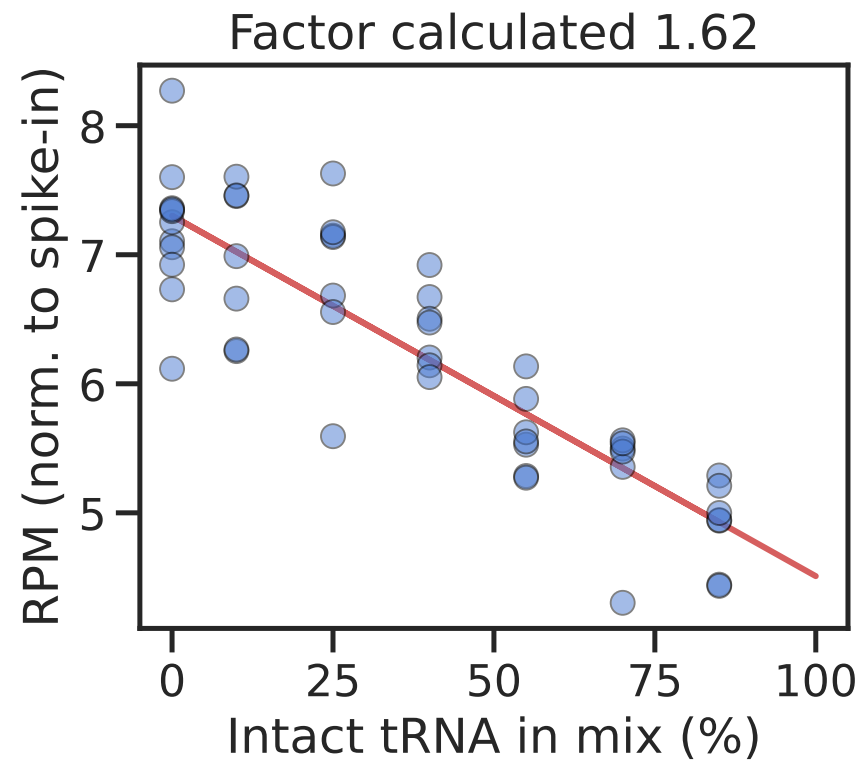
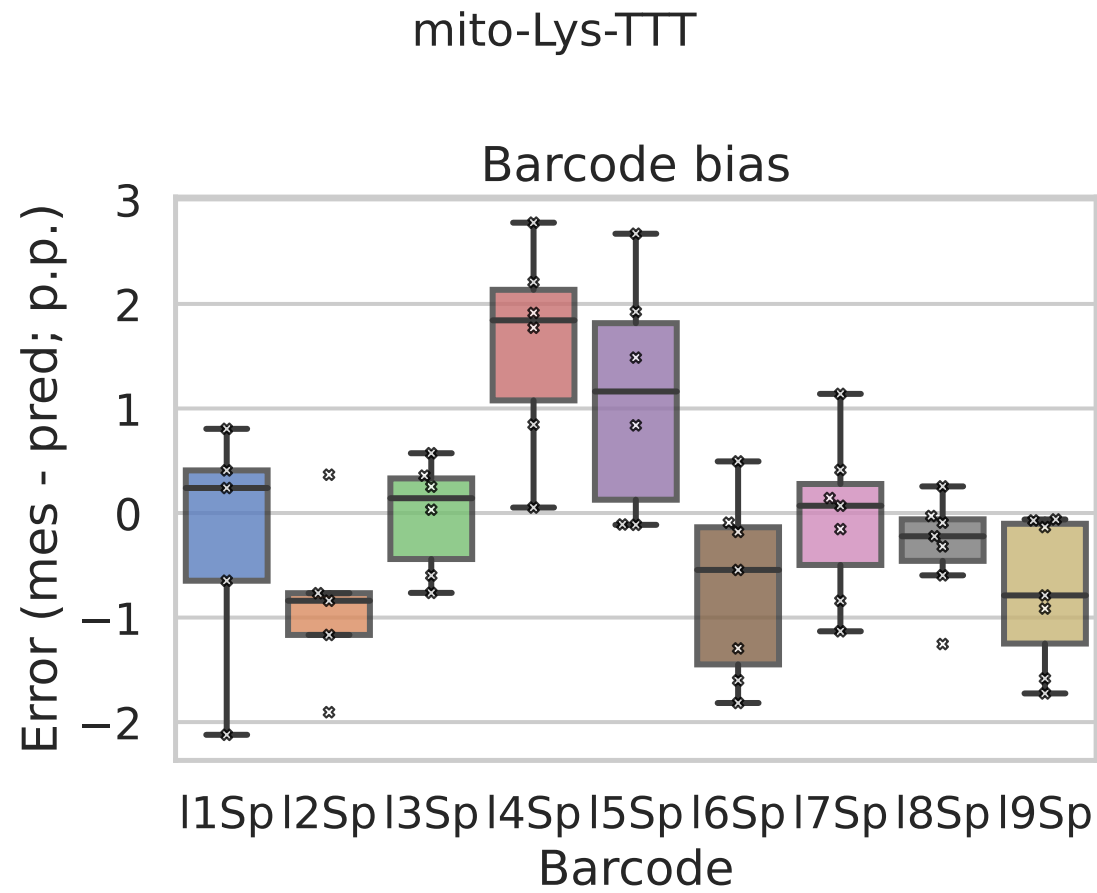
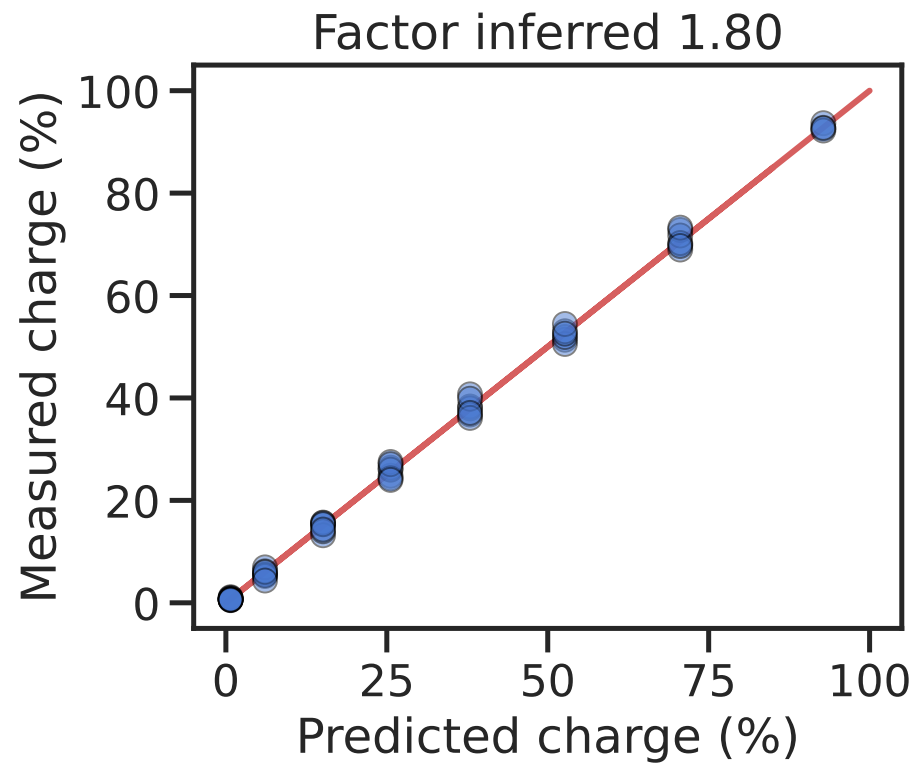


Ser-TGA-1-1

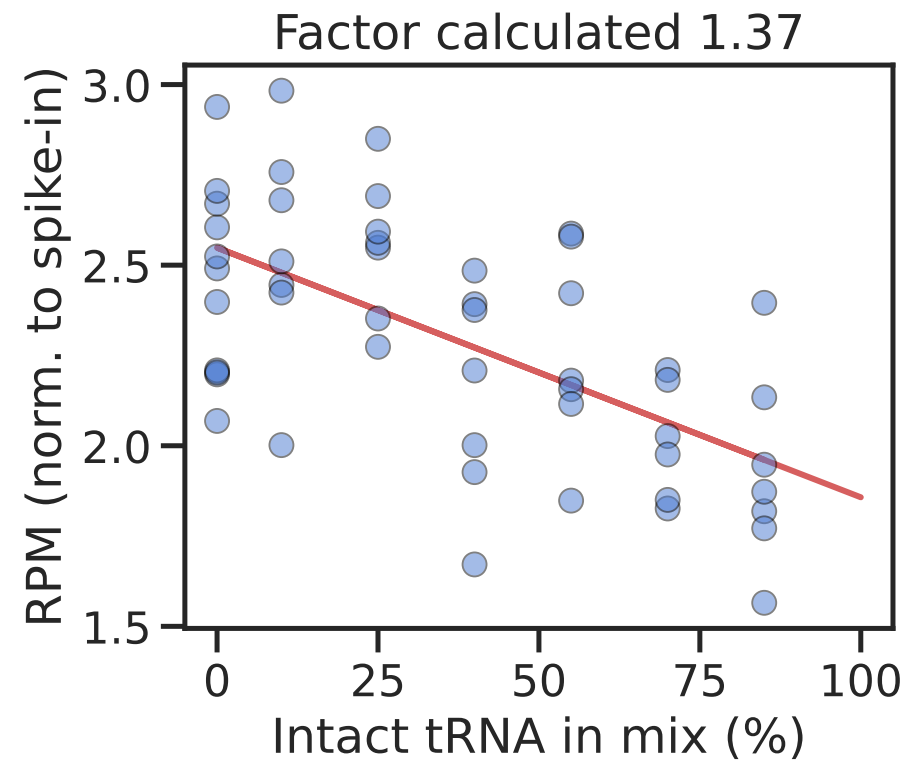
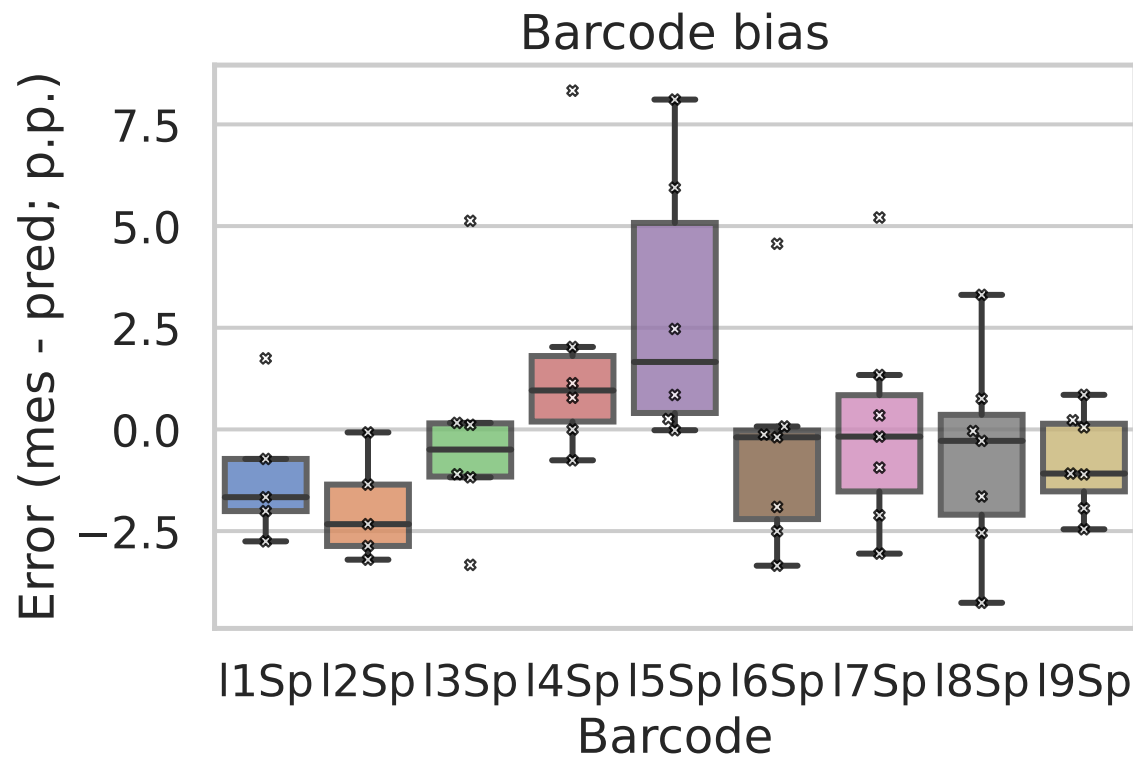
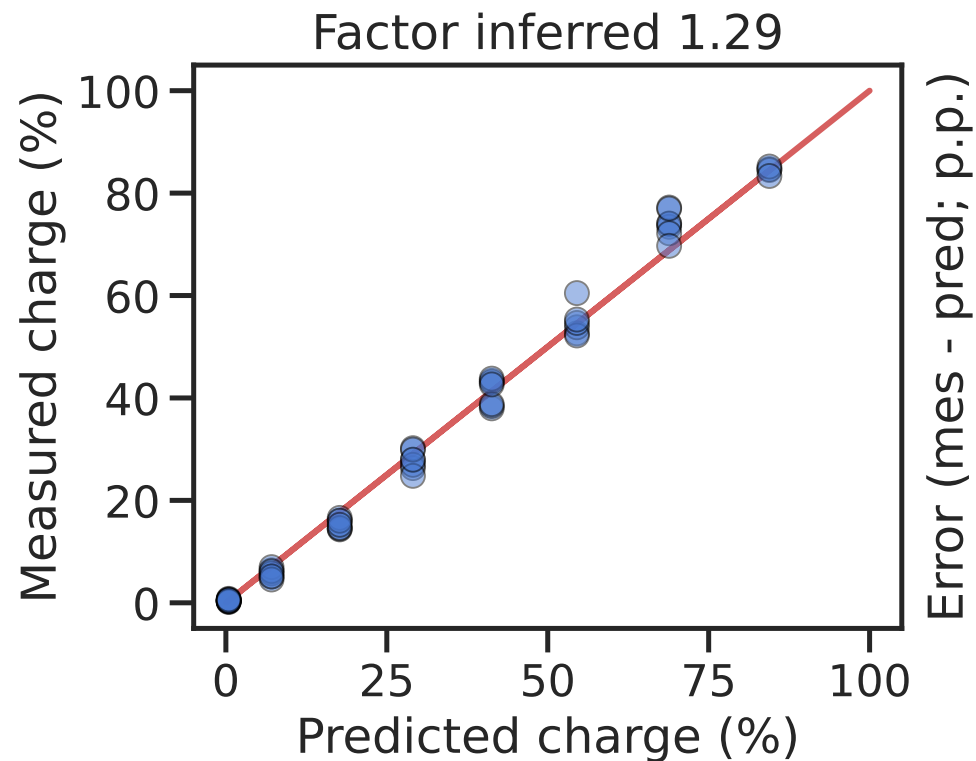


Lys-CTT-4-1

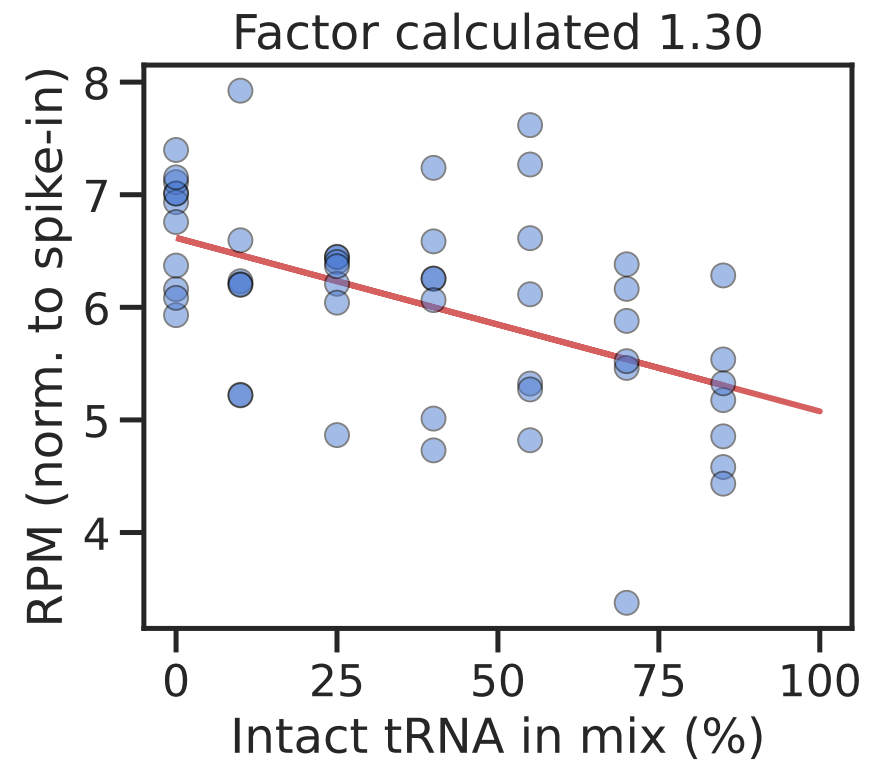
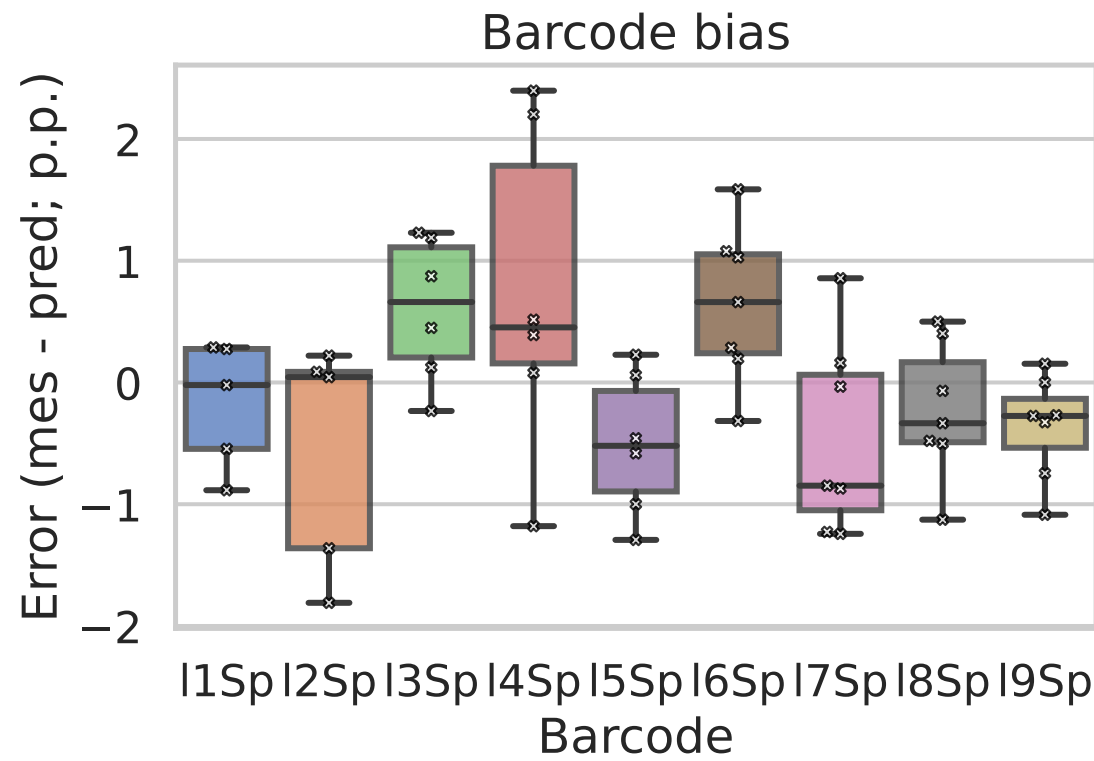
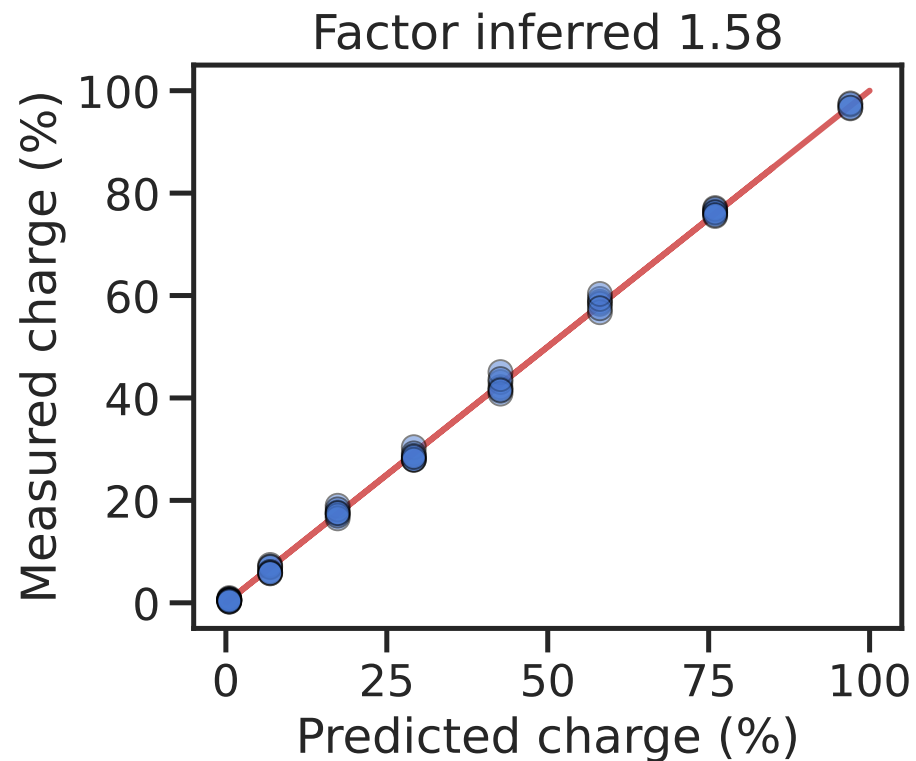




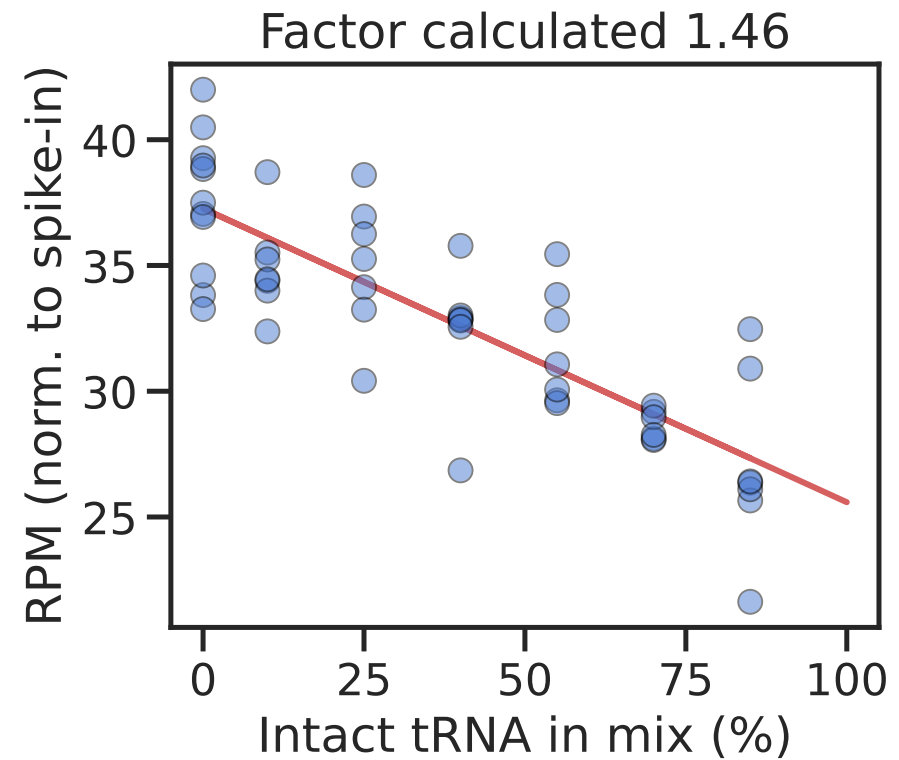
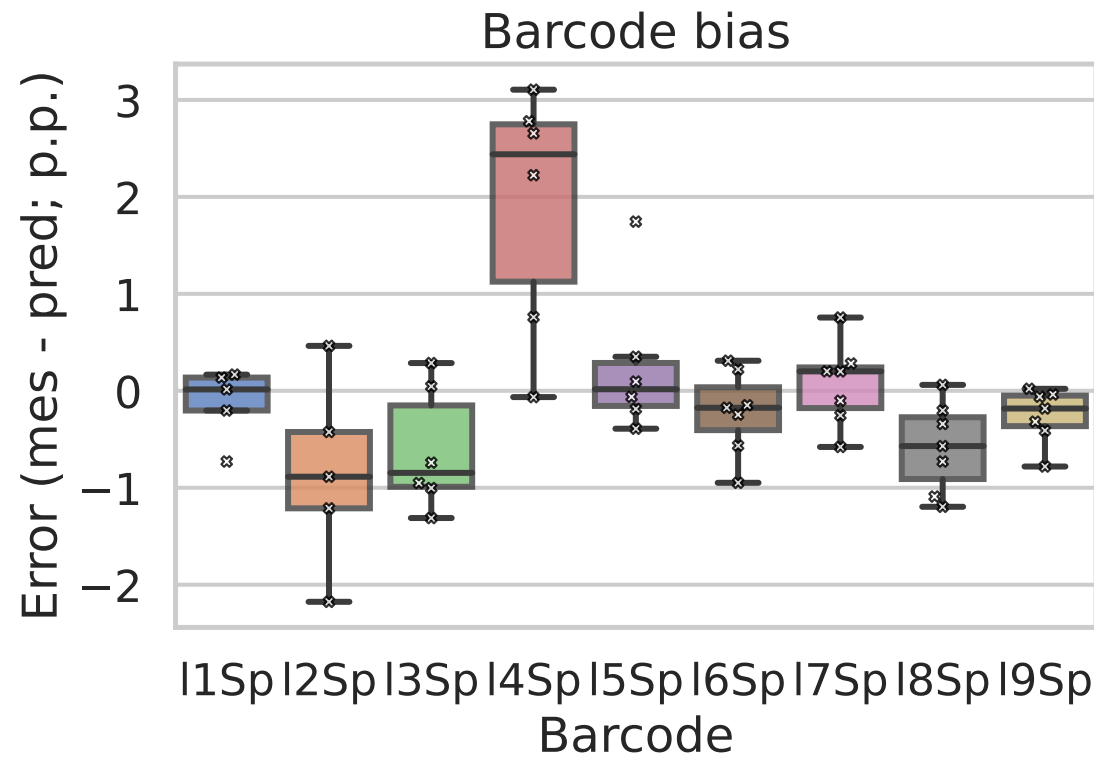
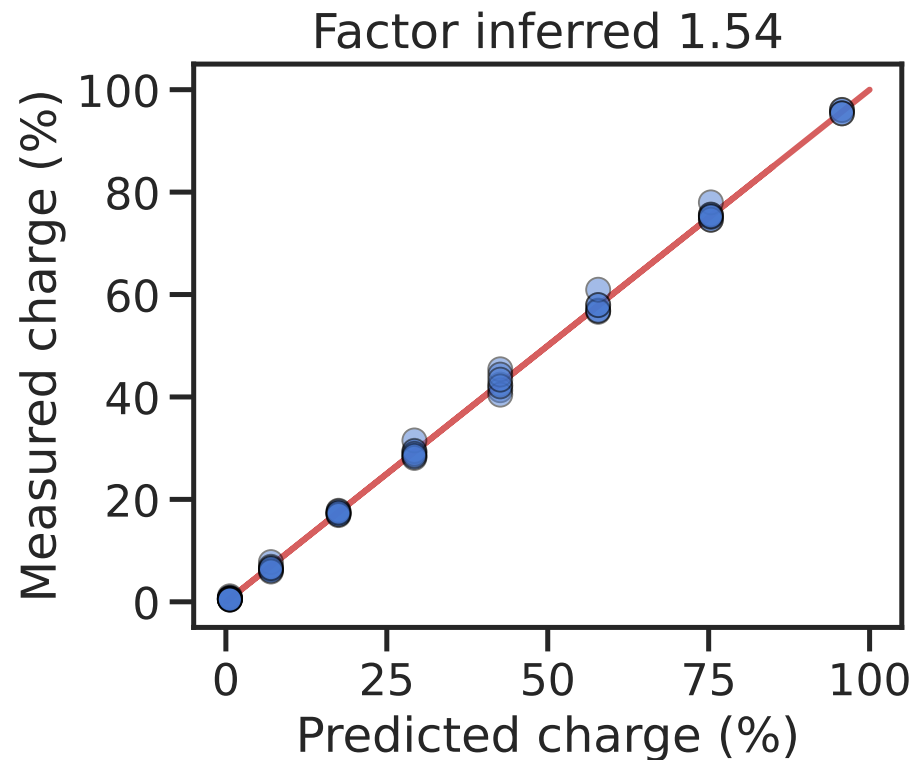
Ser-TGA-3-1



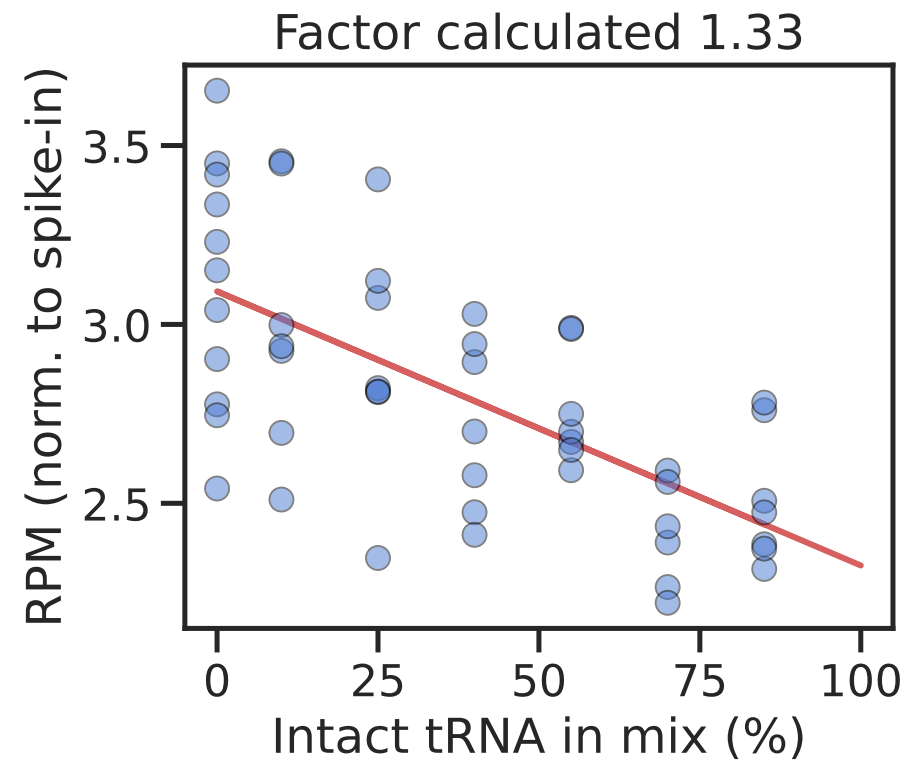
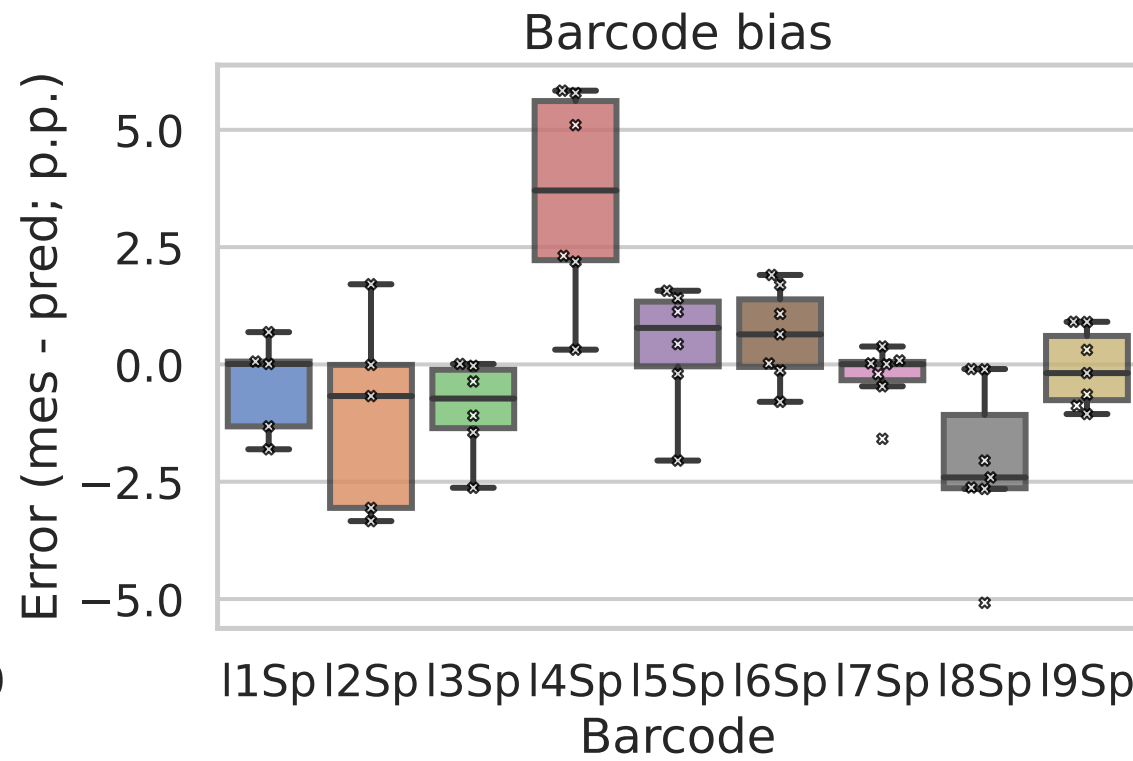
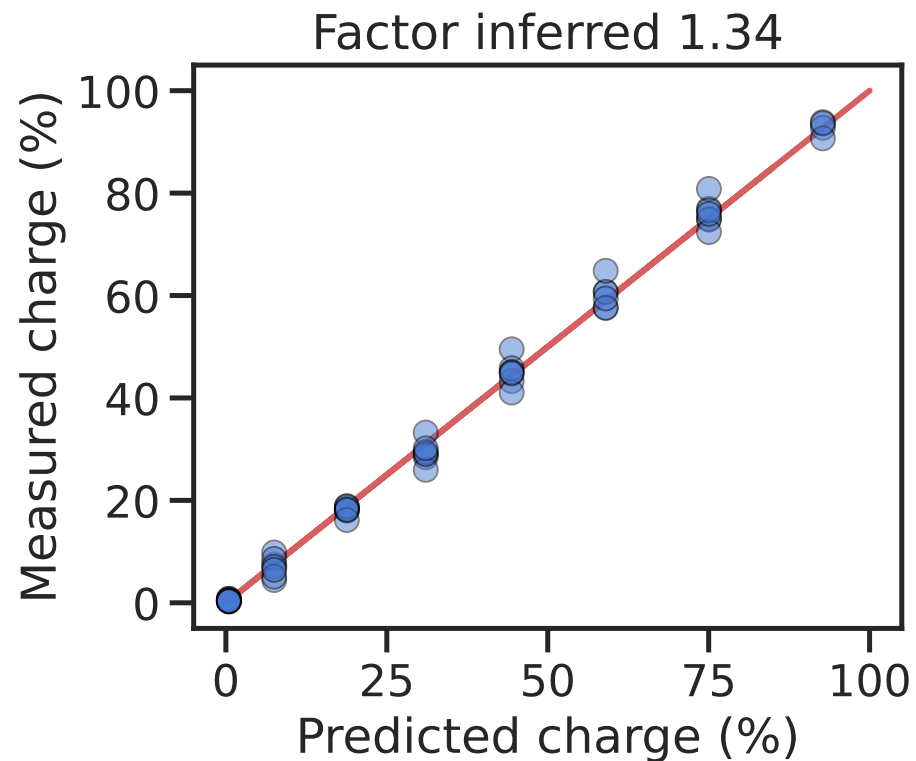
Leu-TAG-1-1



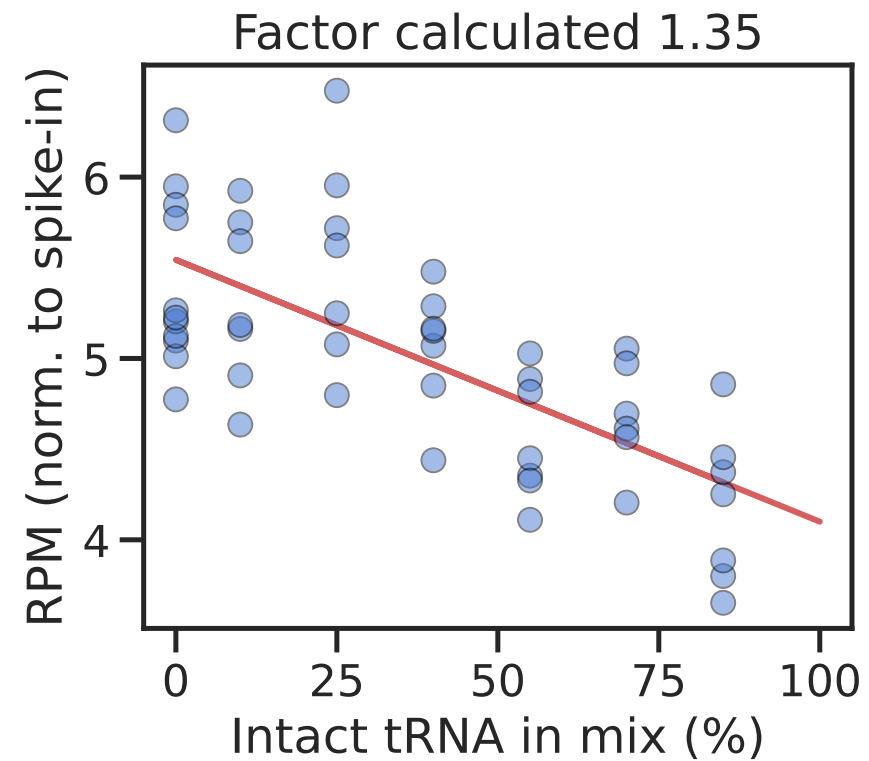
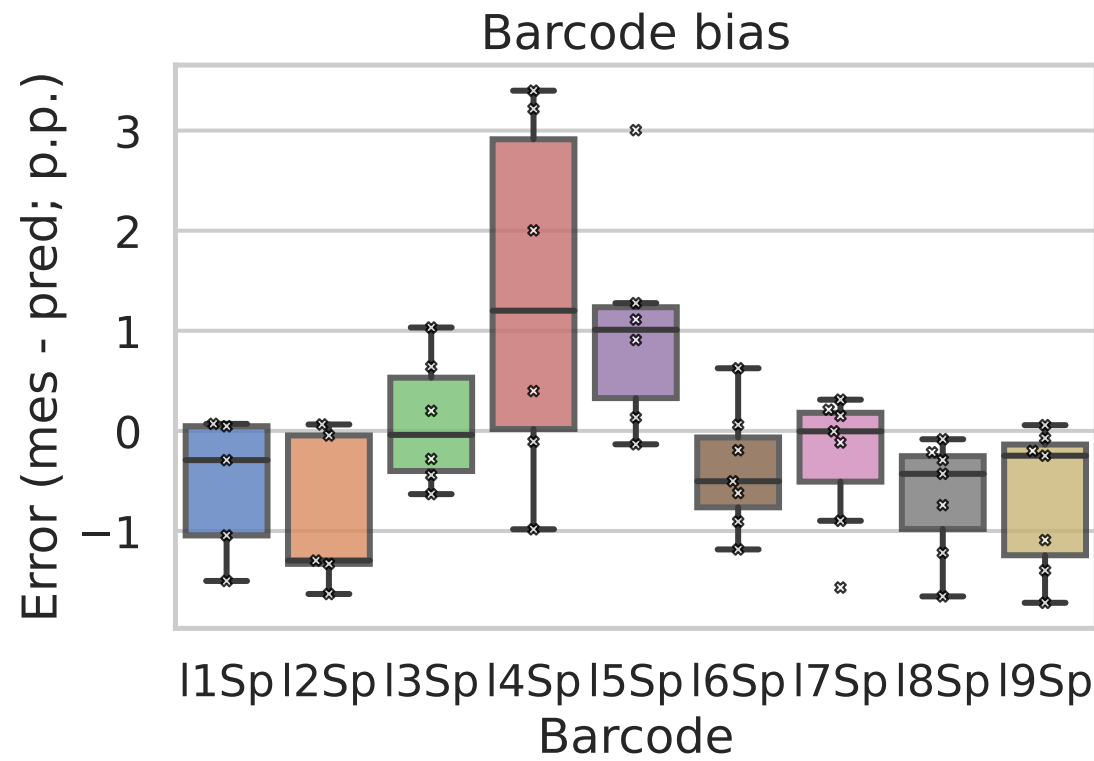
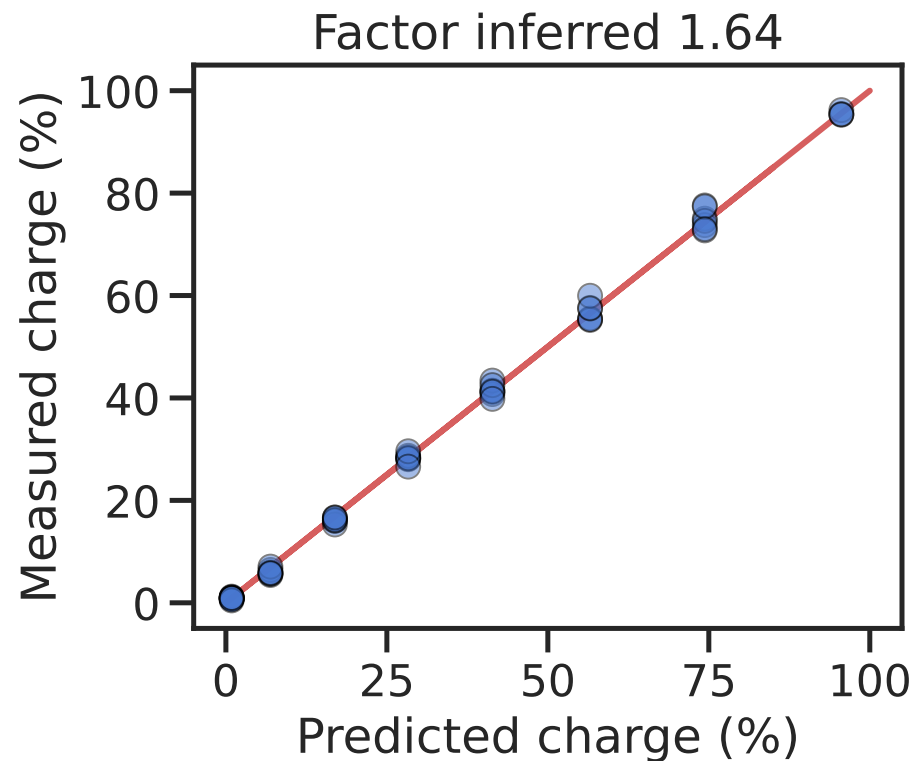
Ala-AGC-8-1



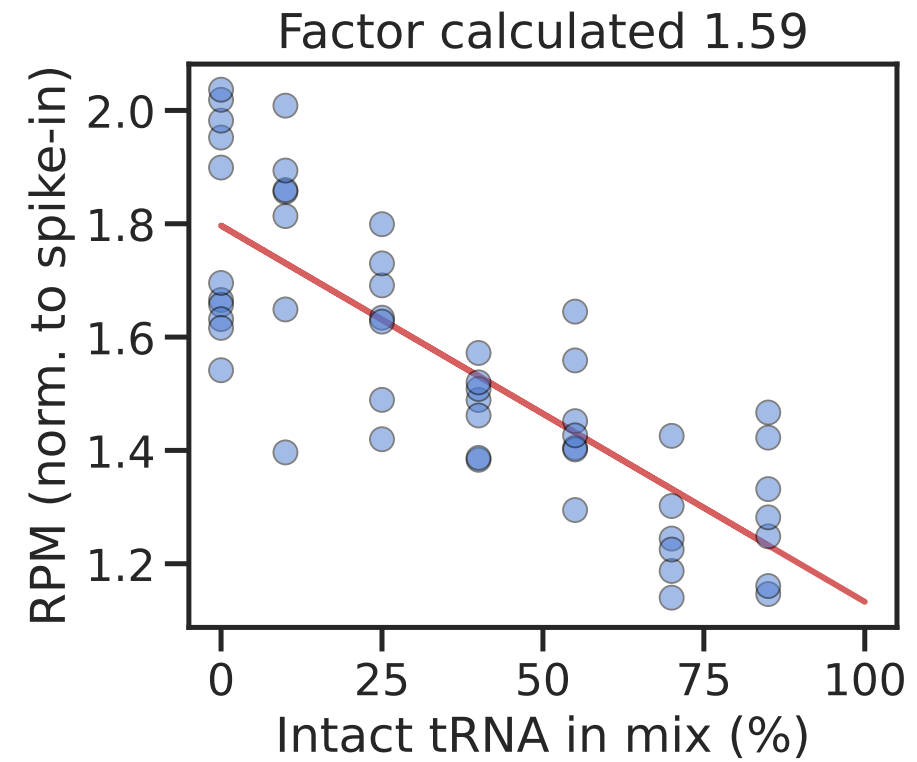
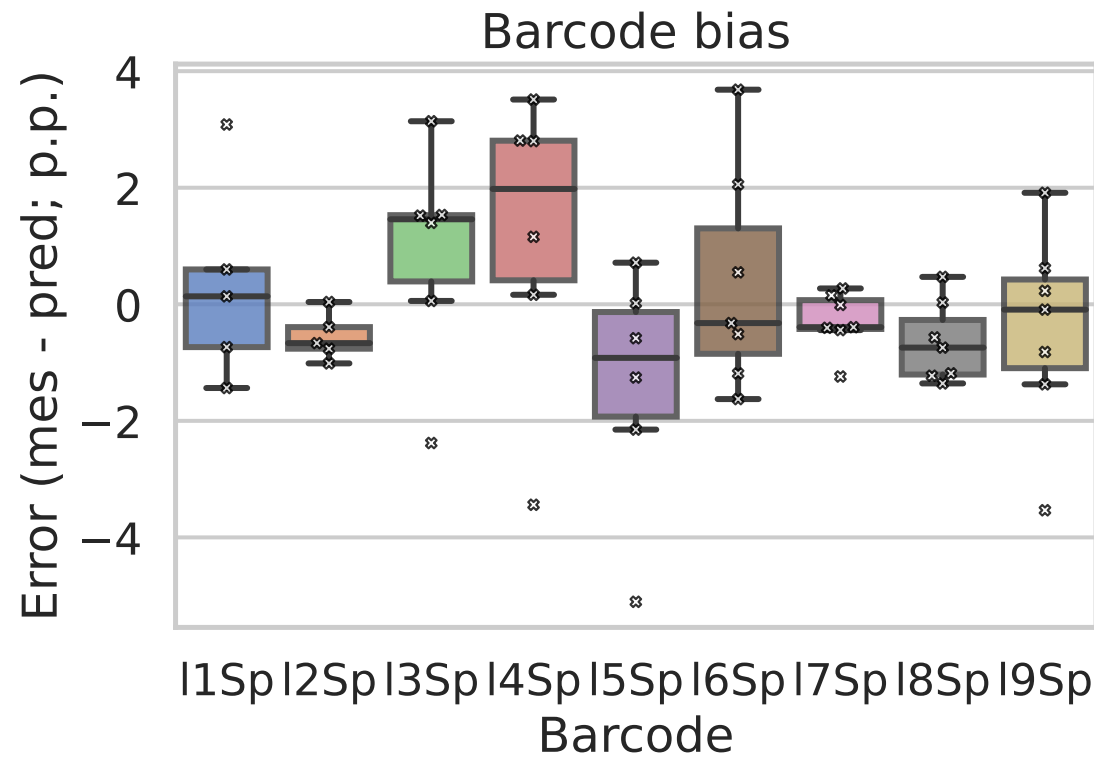
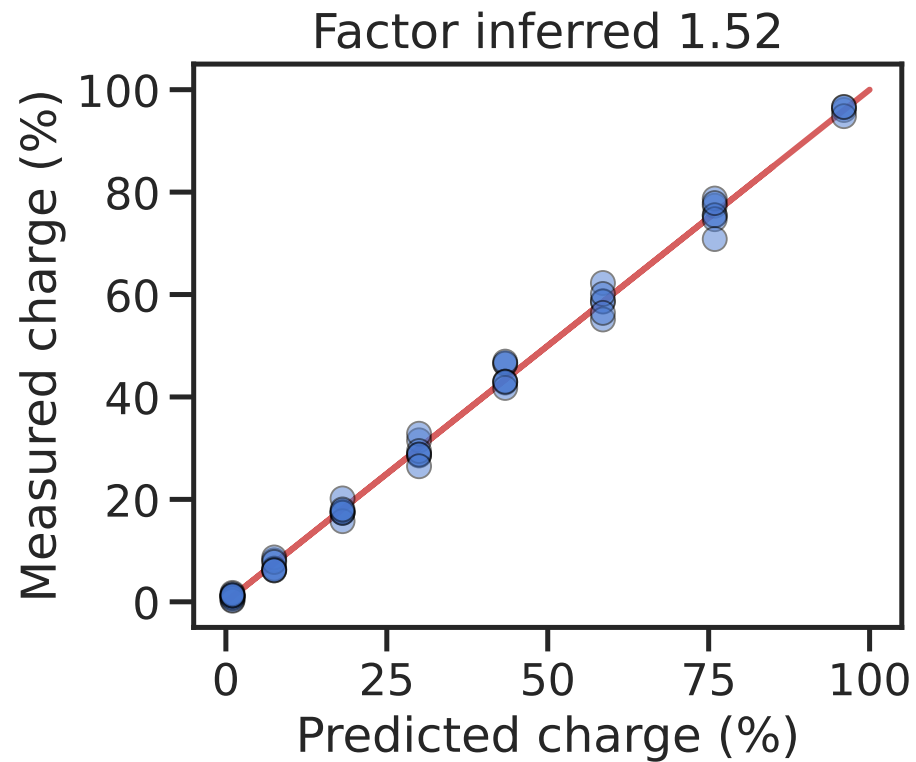
Thr-CGT-3-1



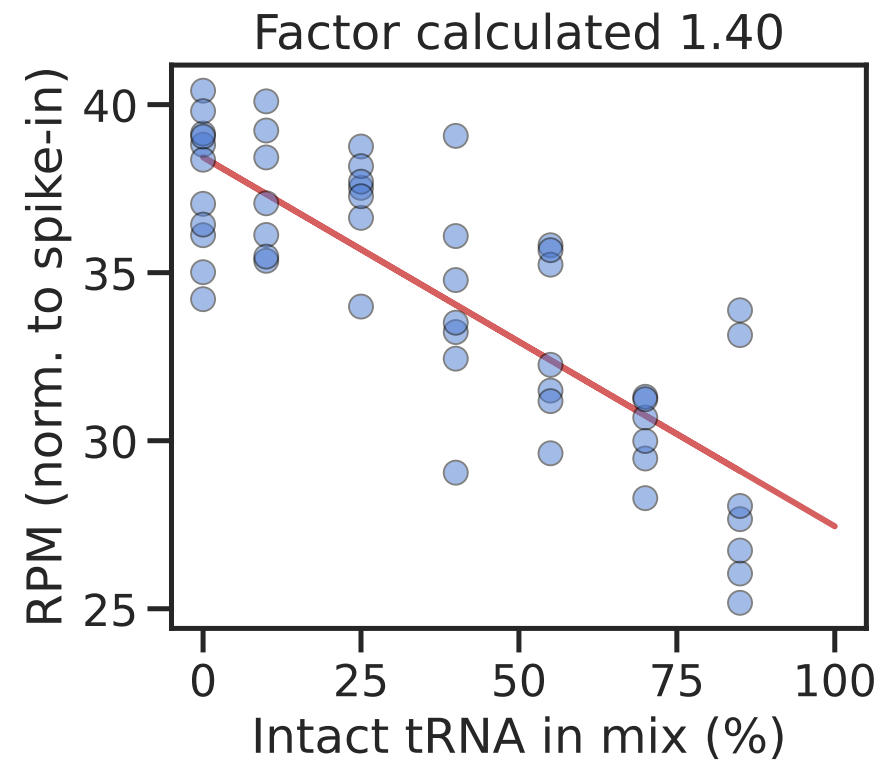
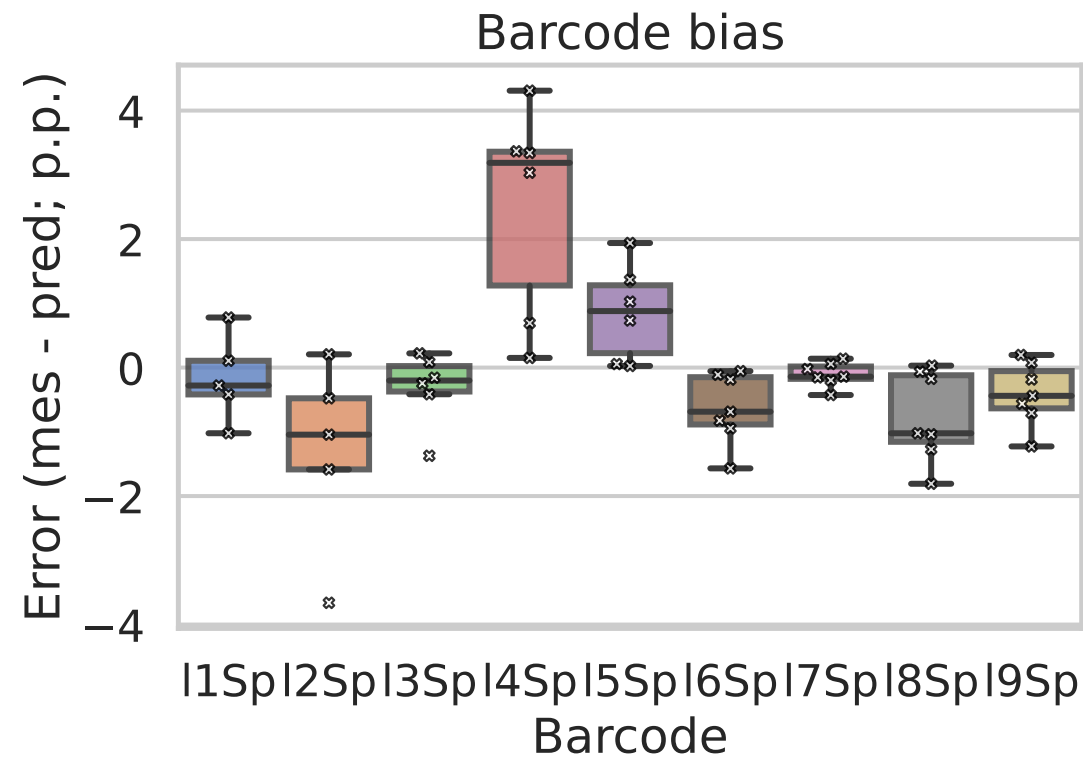
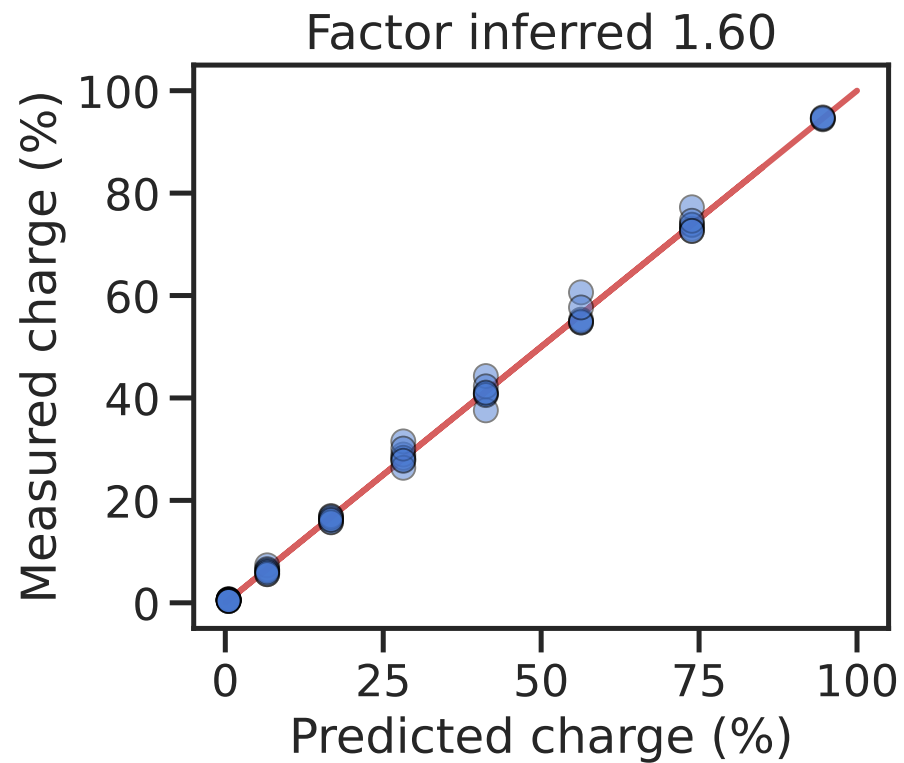
mito-Arg-TCG



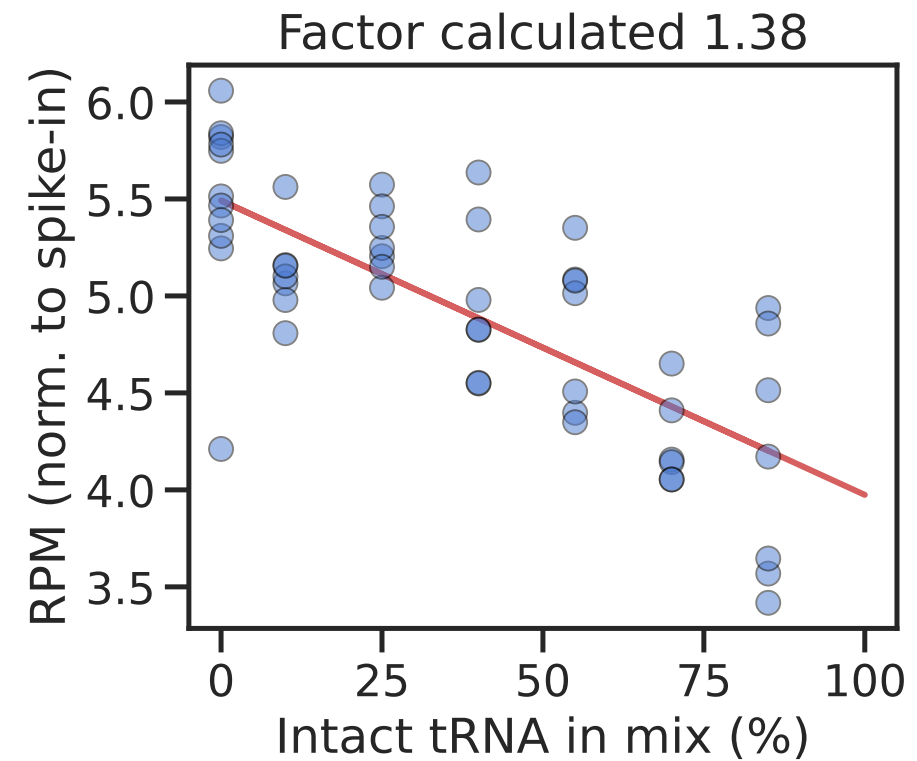
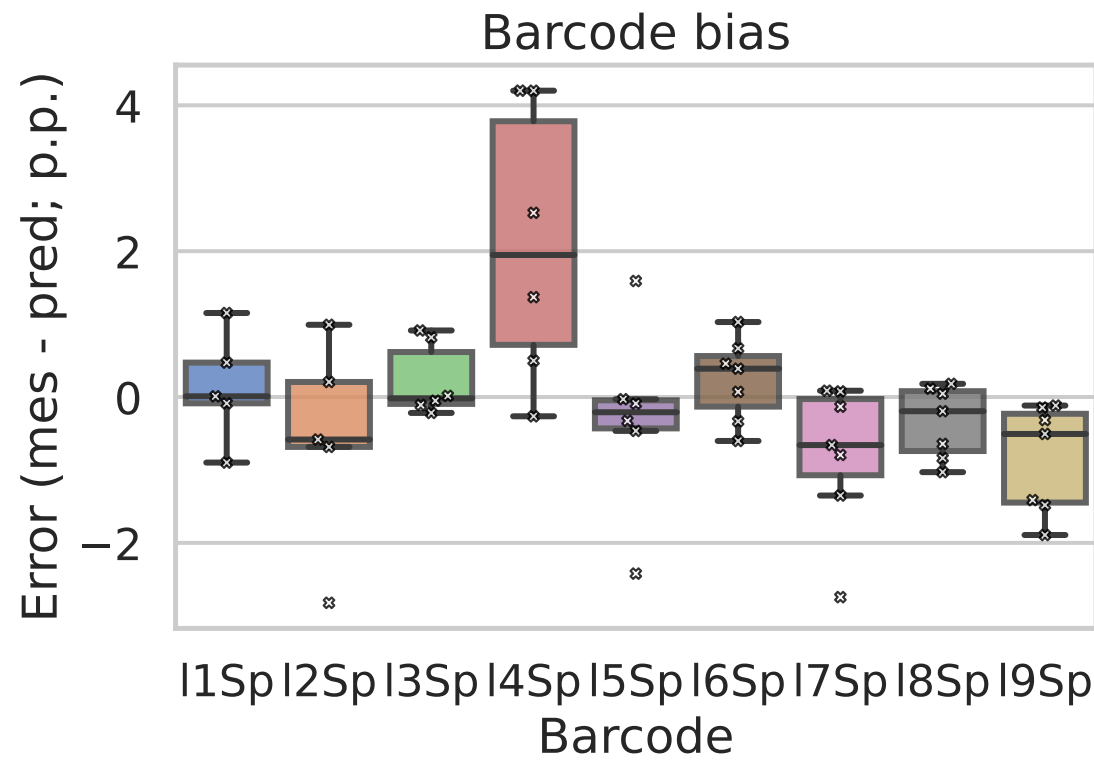
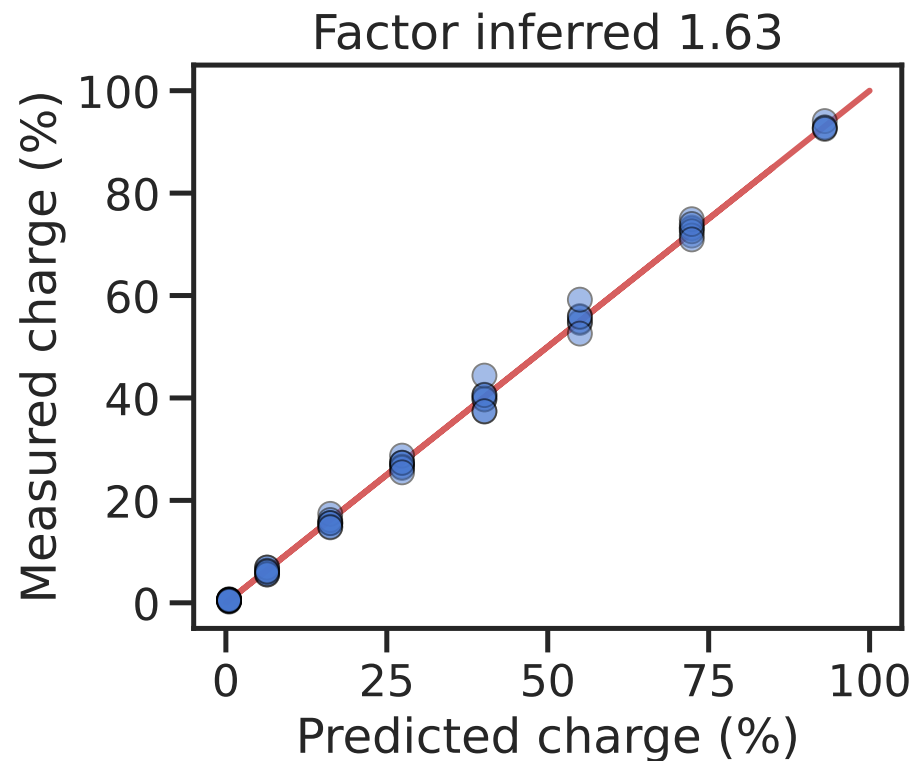
Phe-GAA-3-1



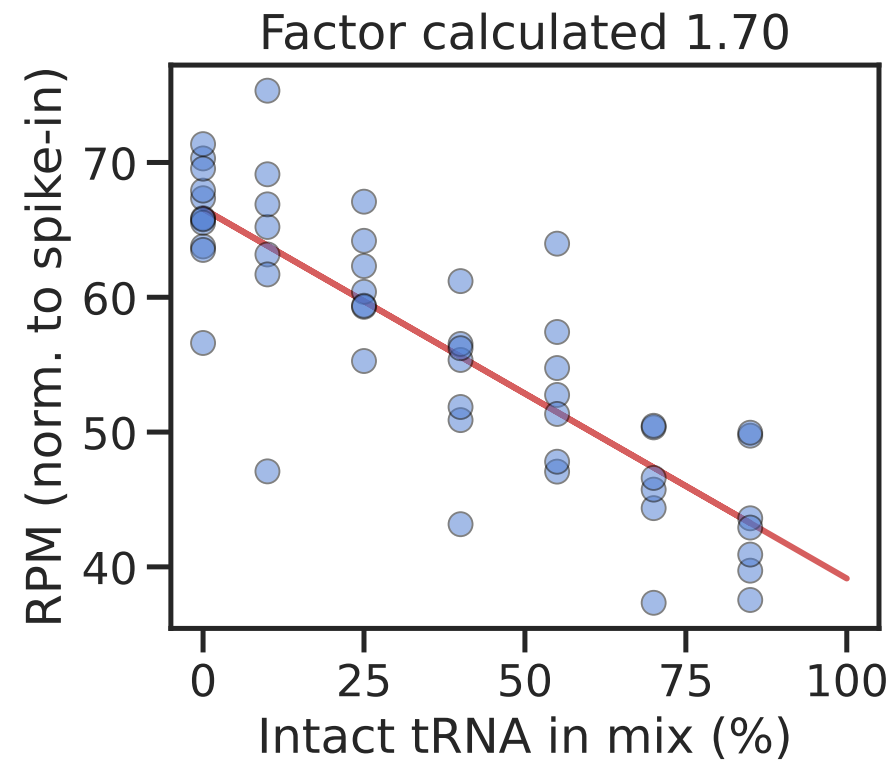
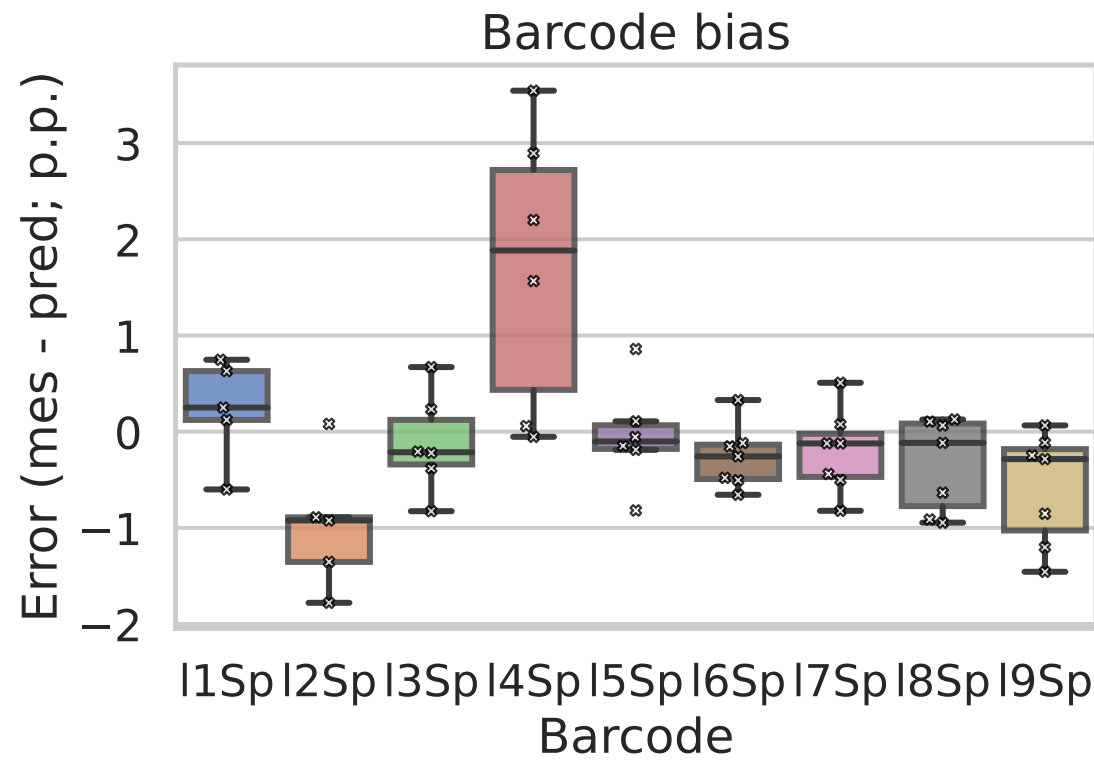
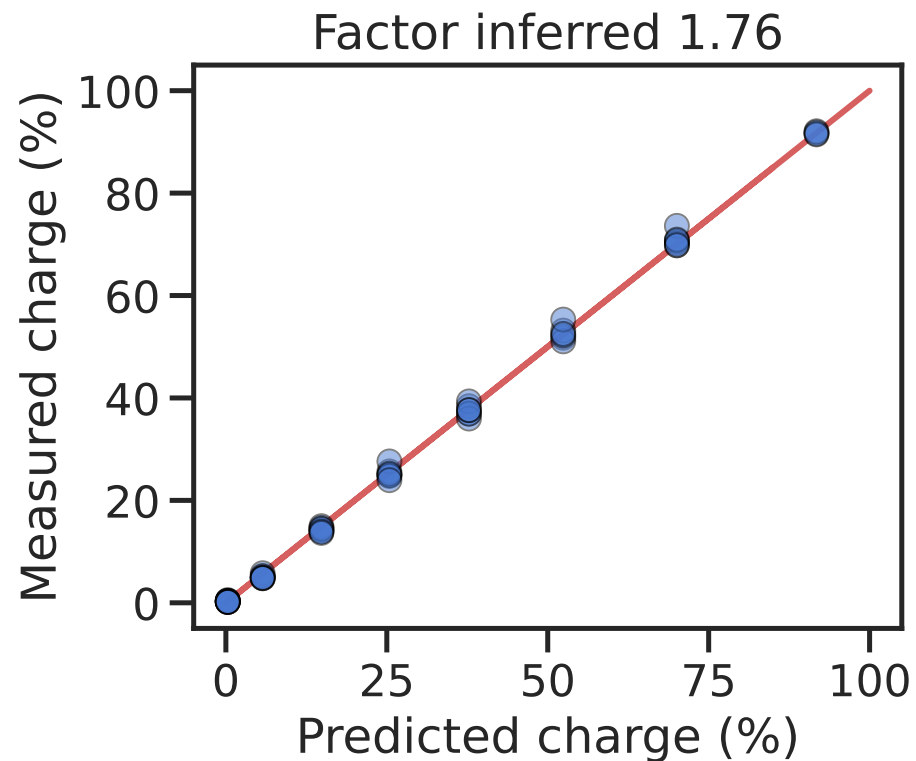
Arg-ACG-1-3



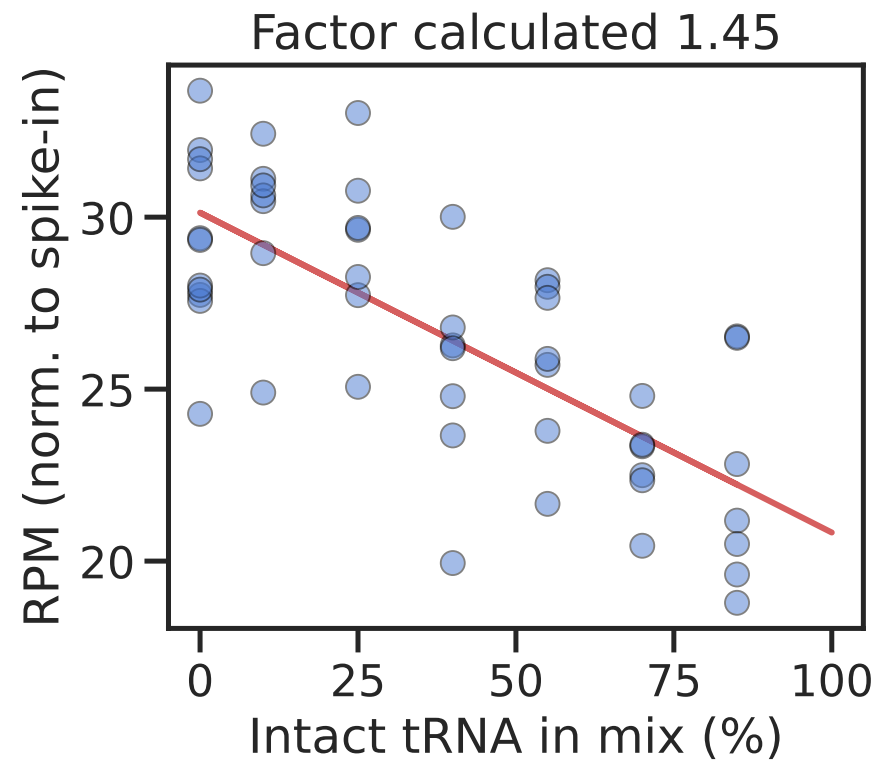
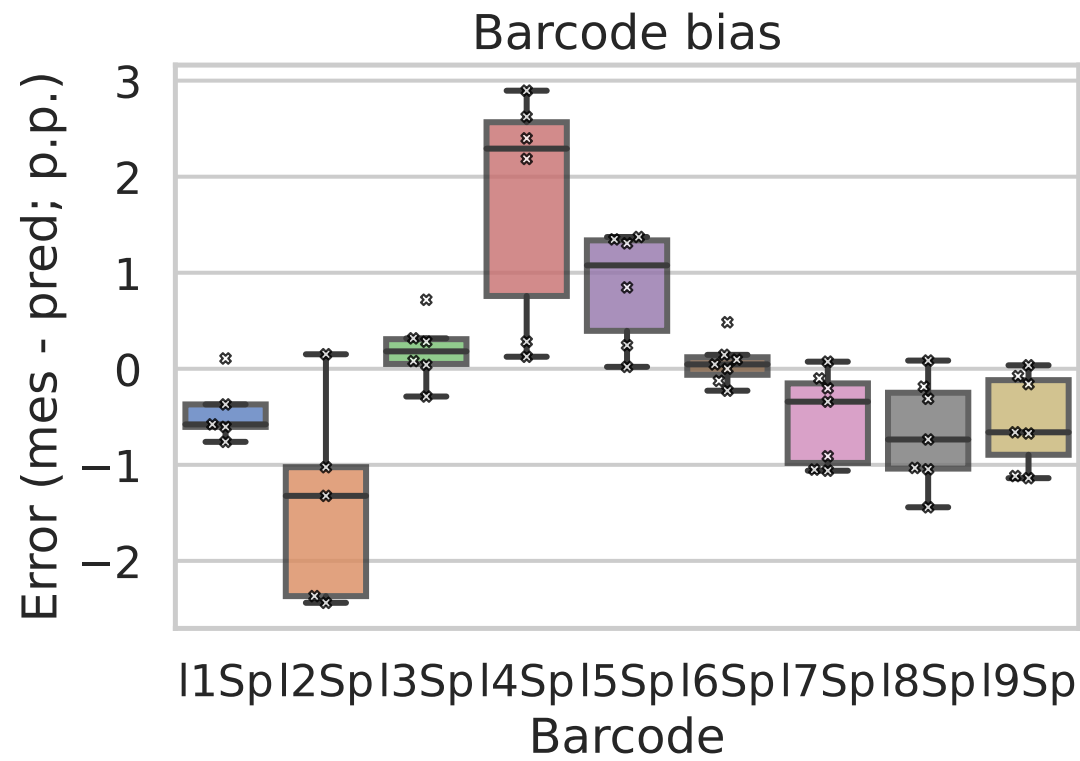
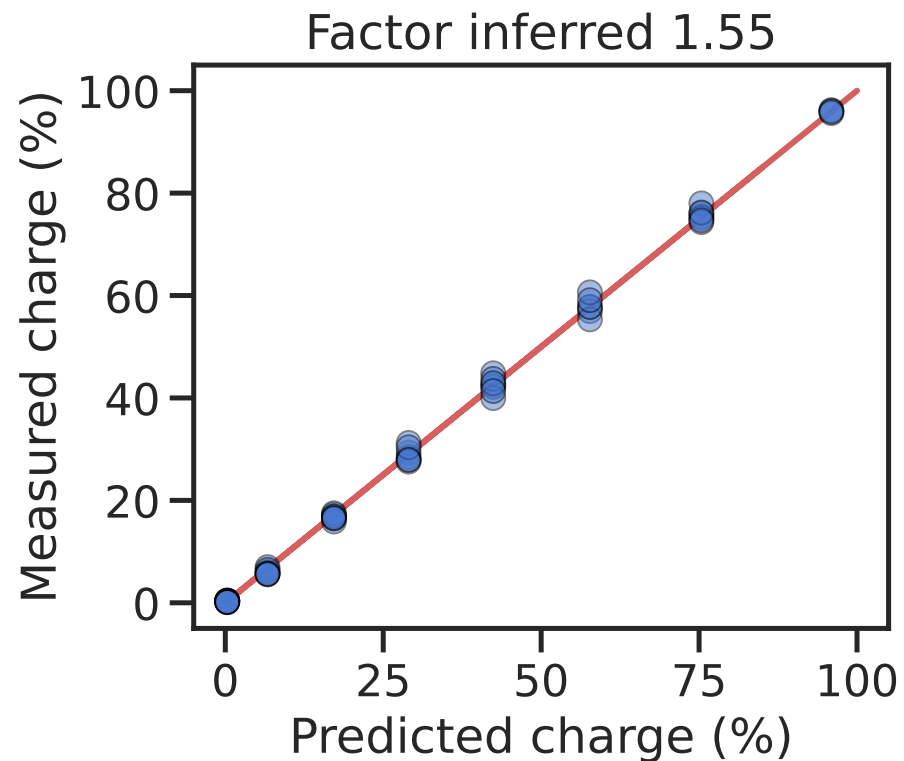
Ala-AGC-6-1



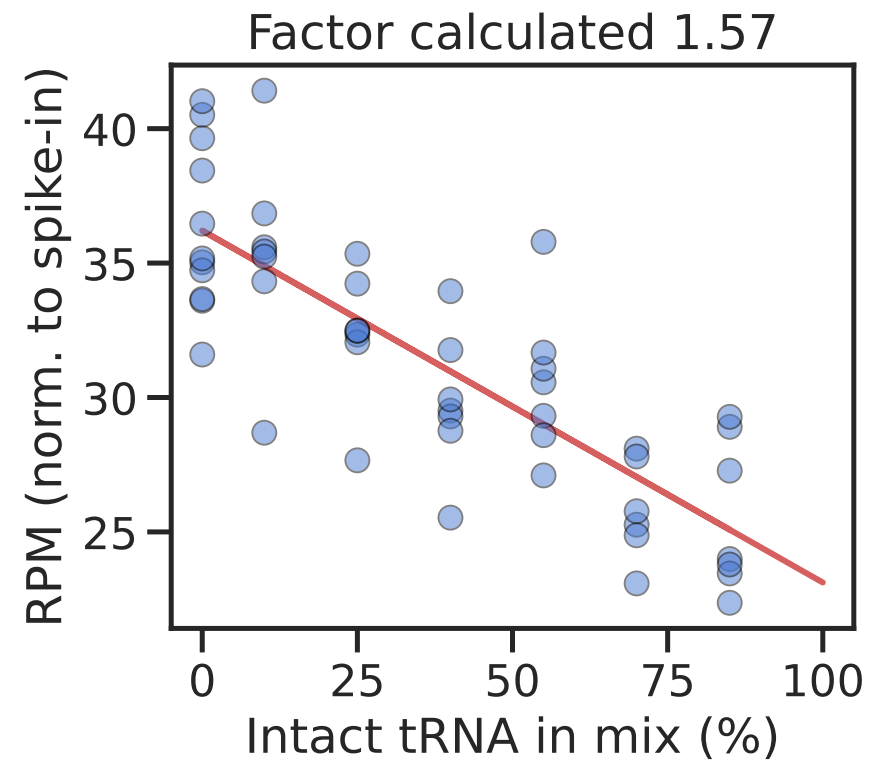
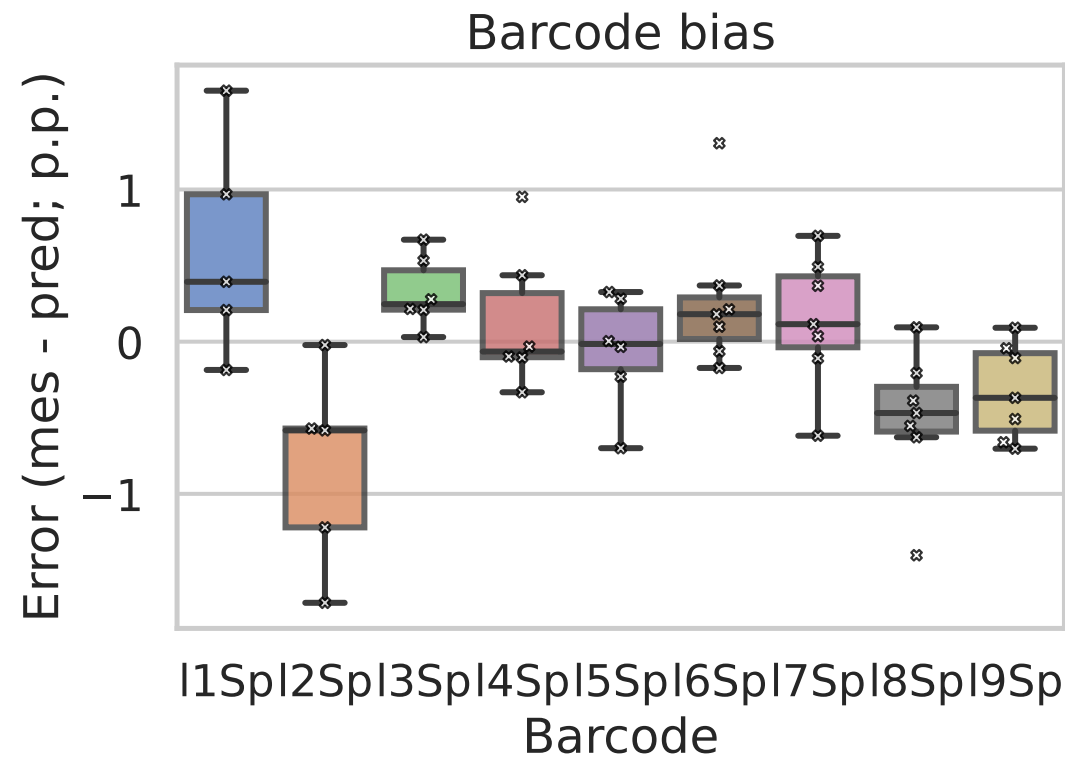
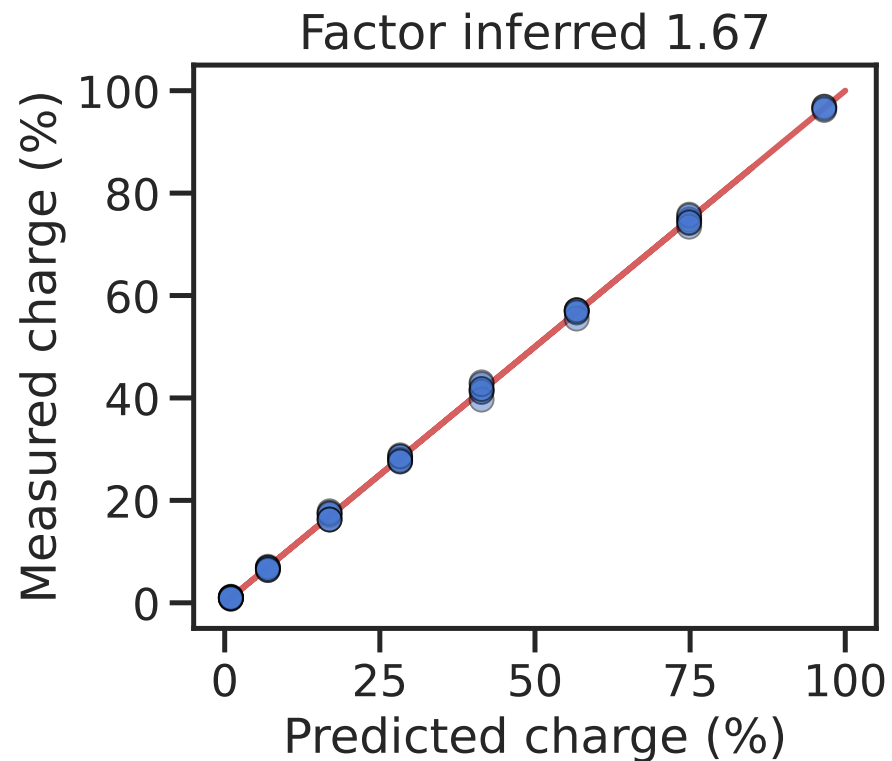
Pro-AGG-1-1



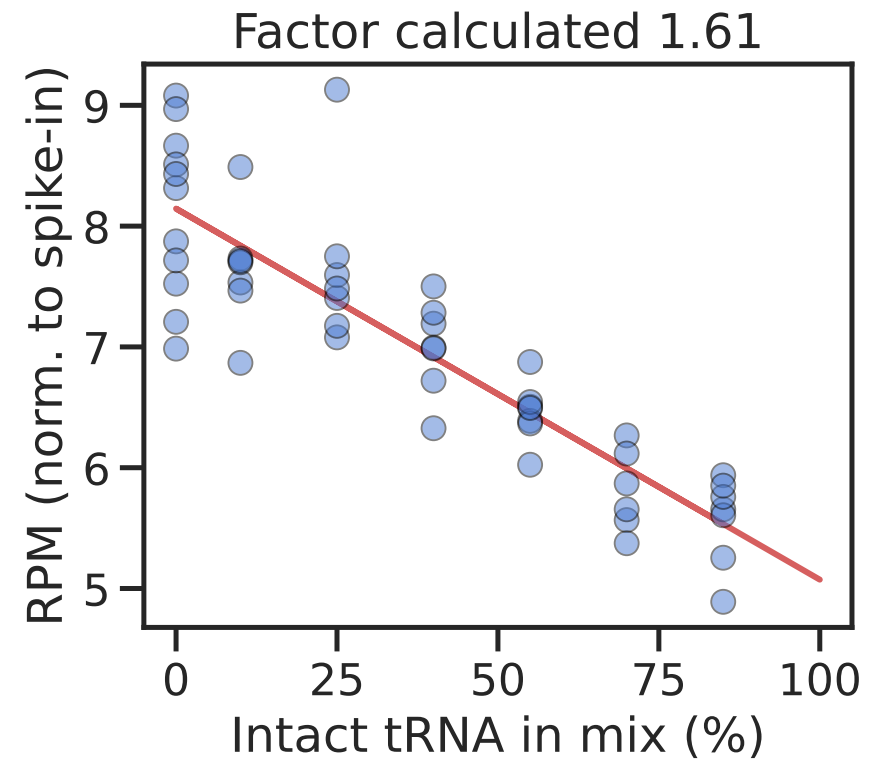
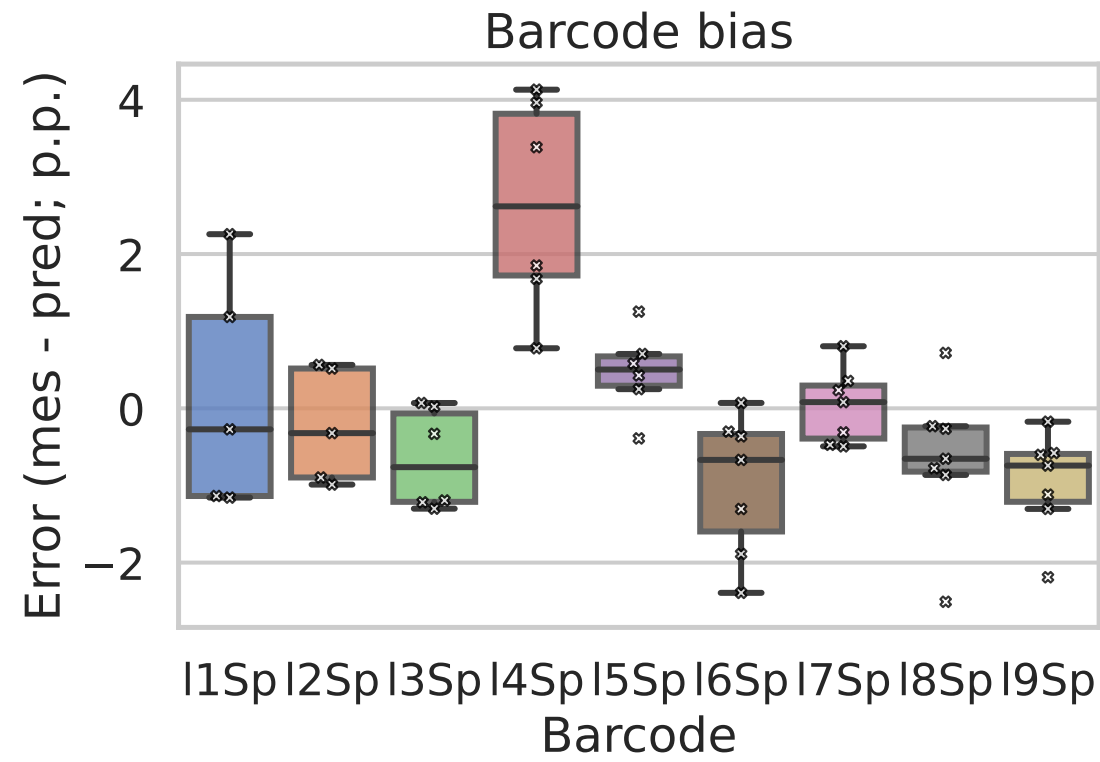
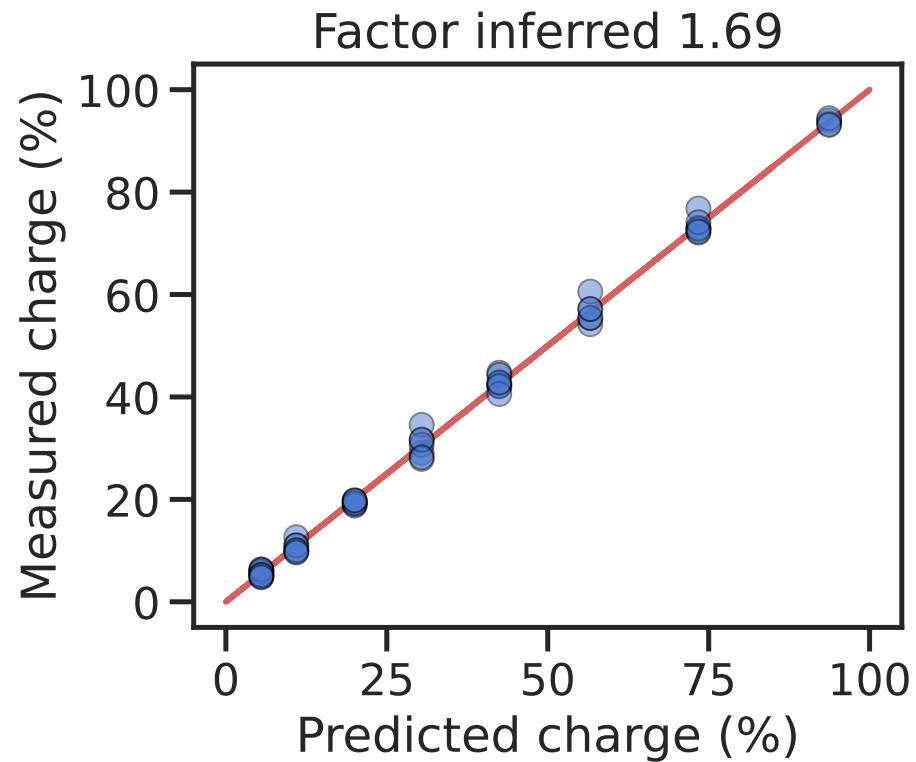
Asn-GTT-2-8



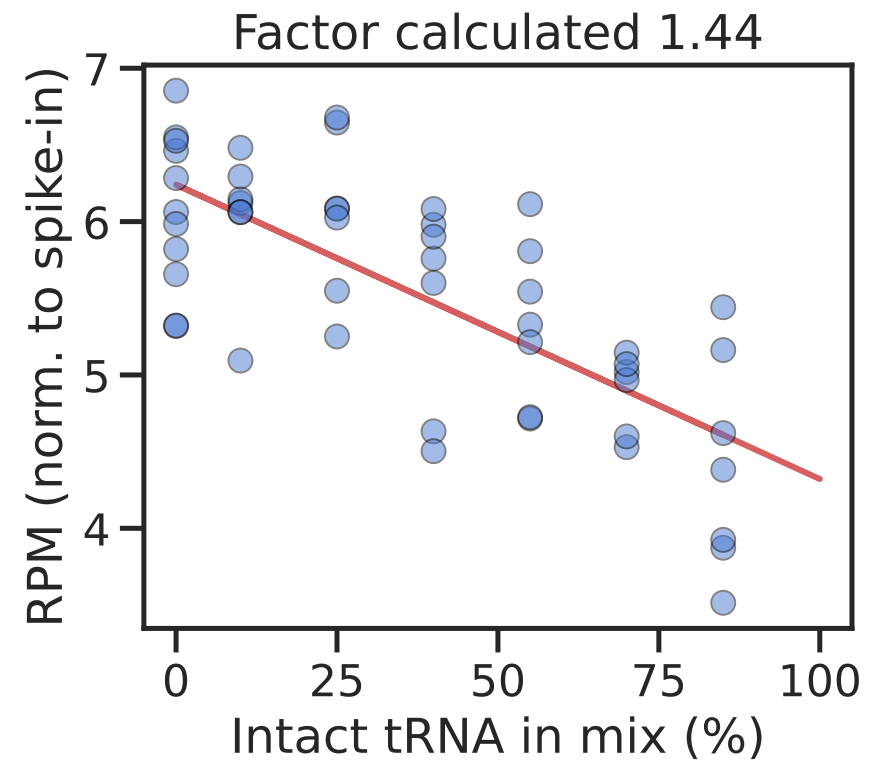
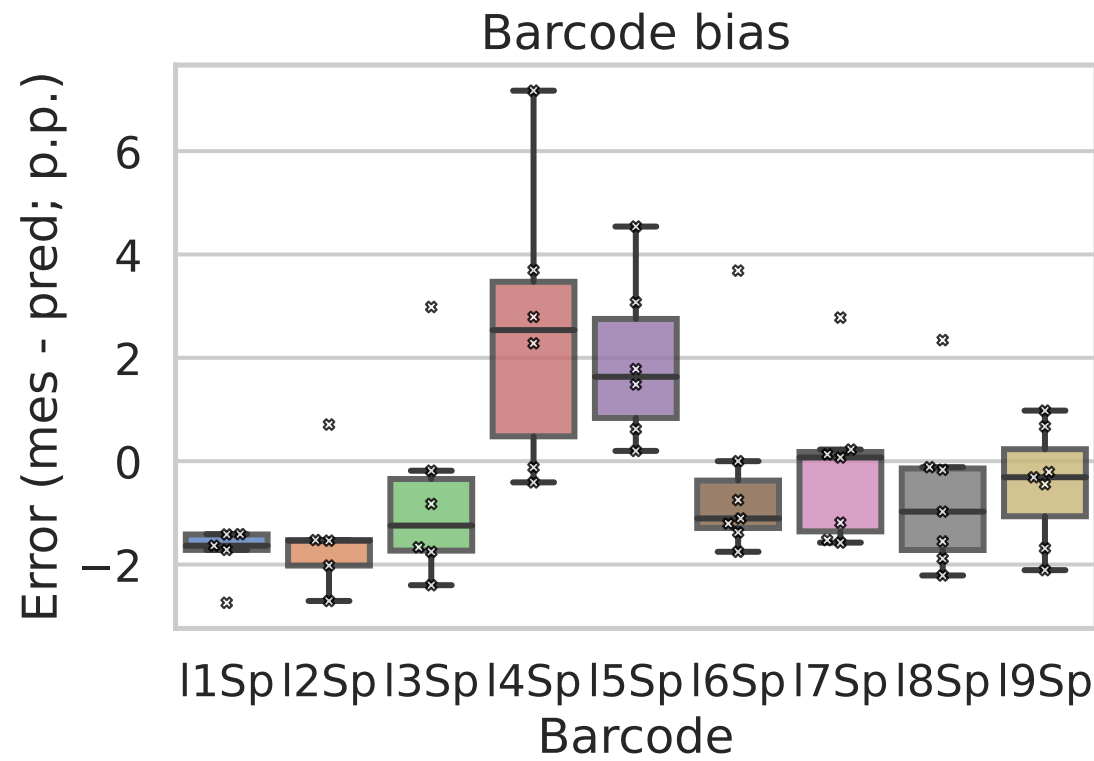
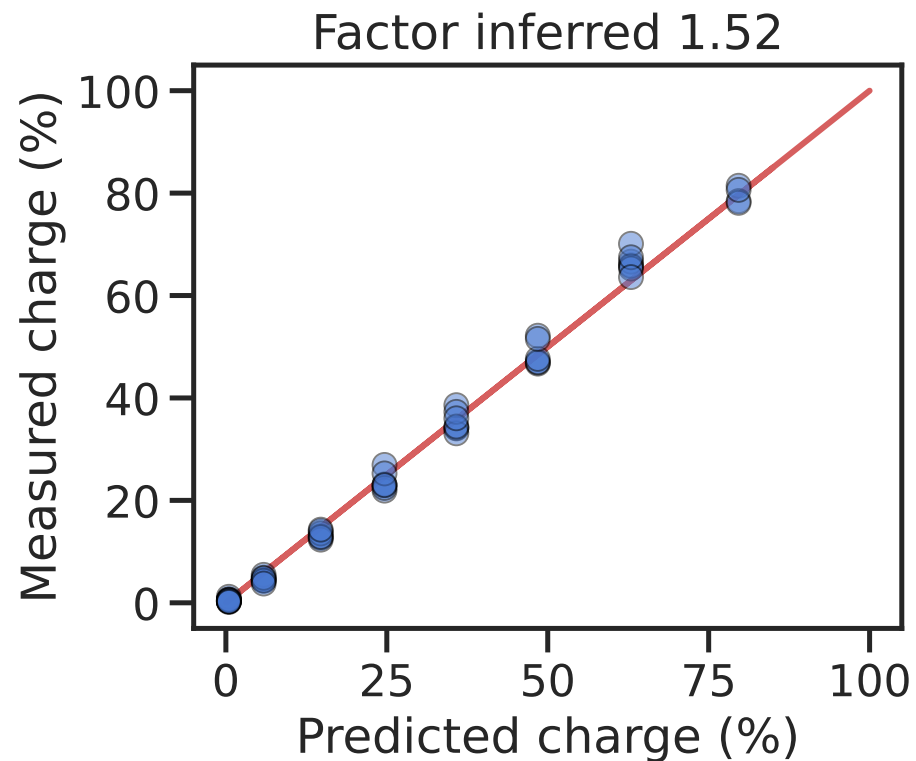
Ile-AAT-5-4



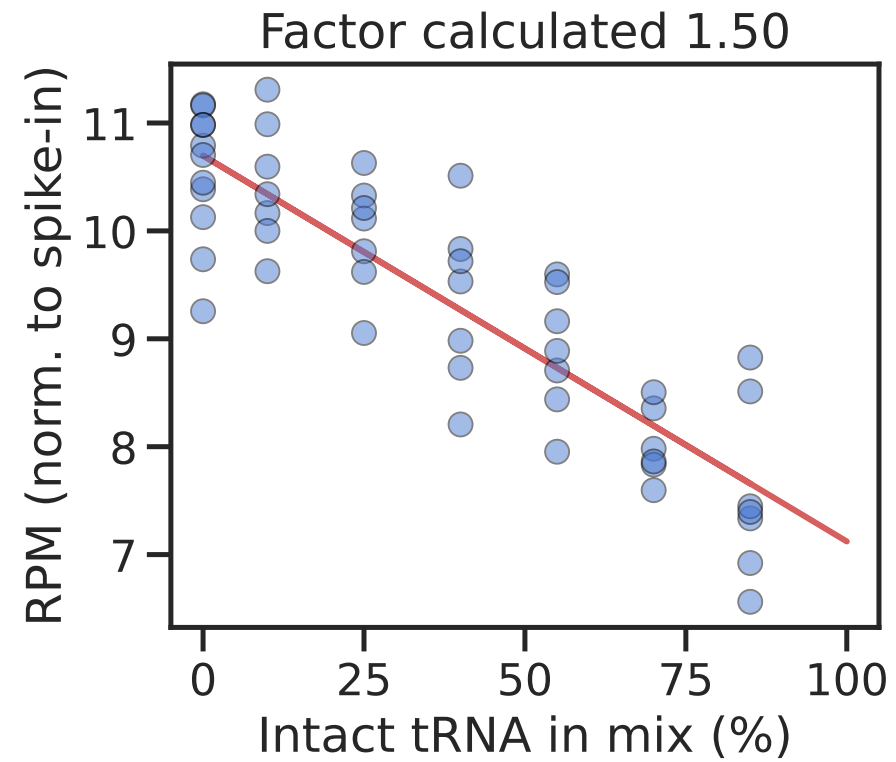
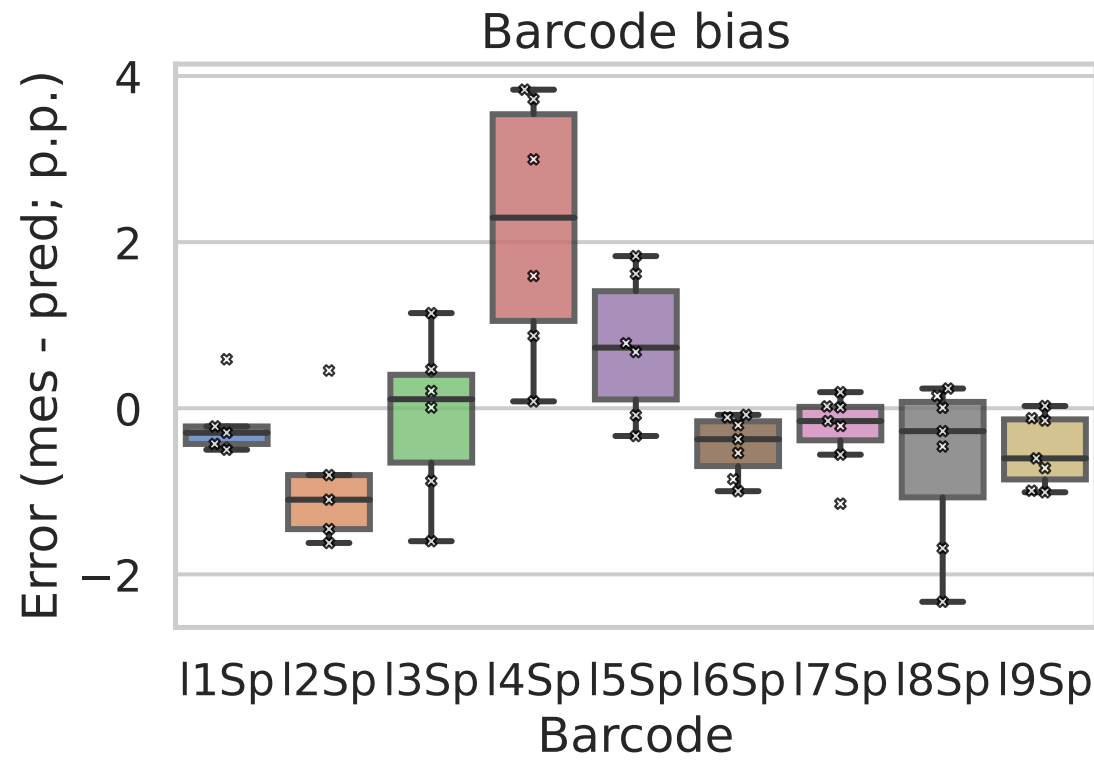
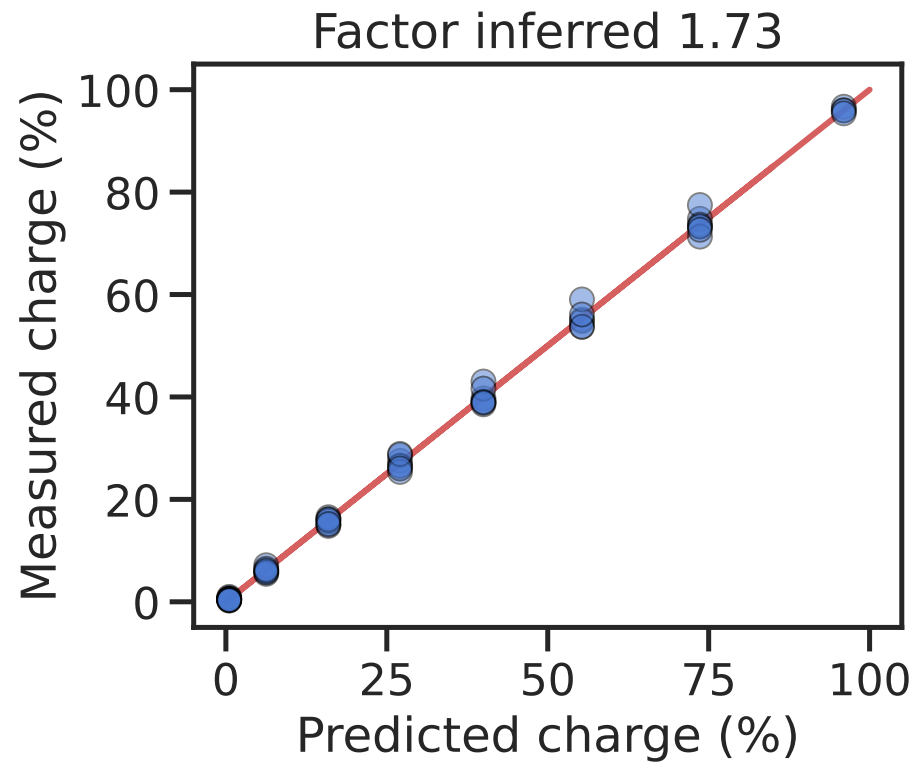
mito-Met-CAT



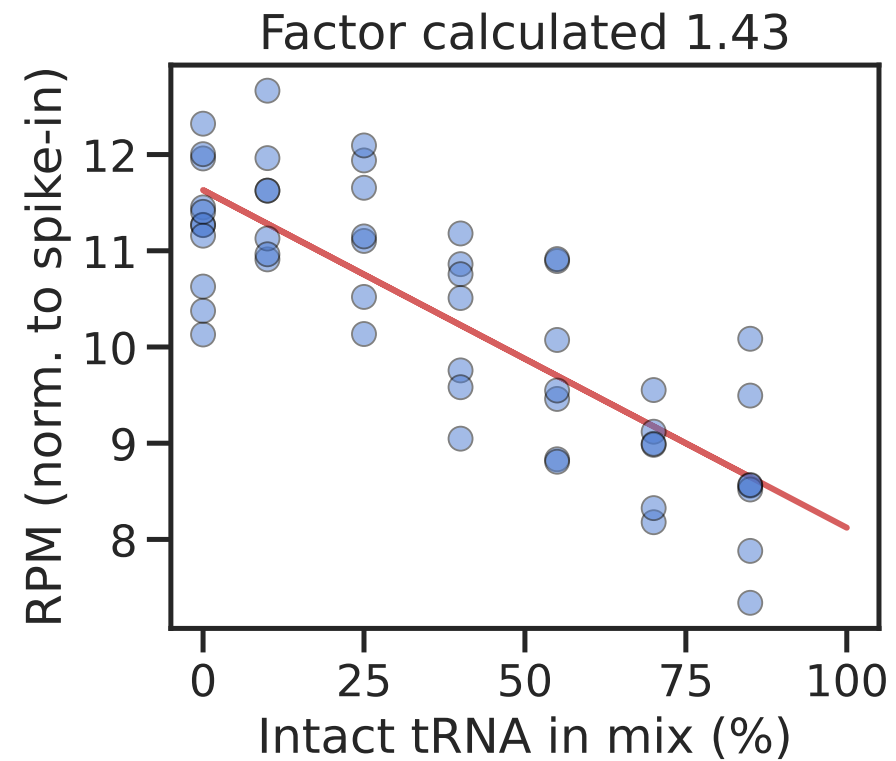
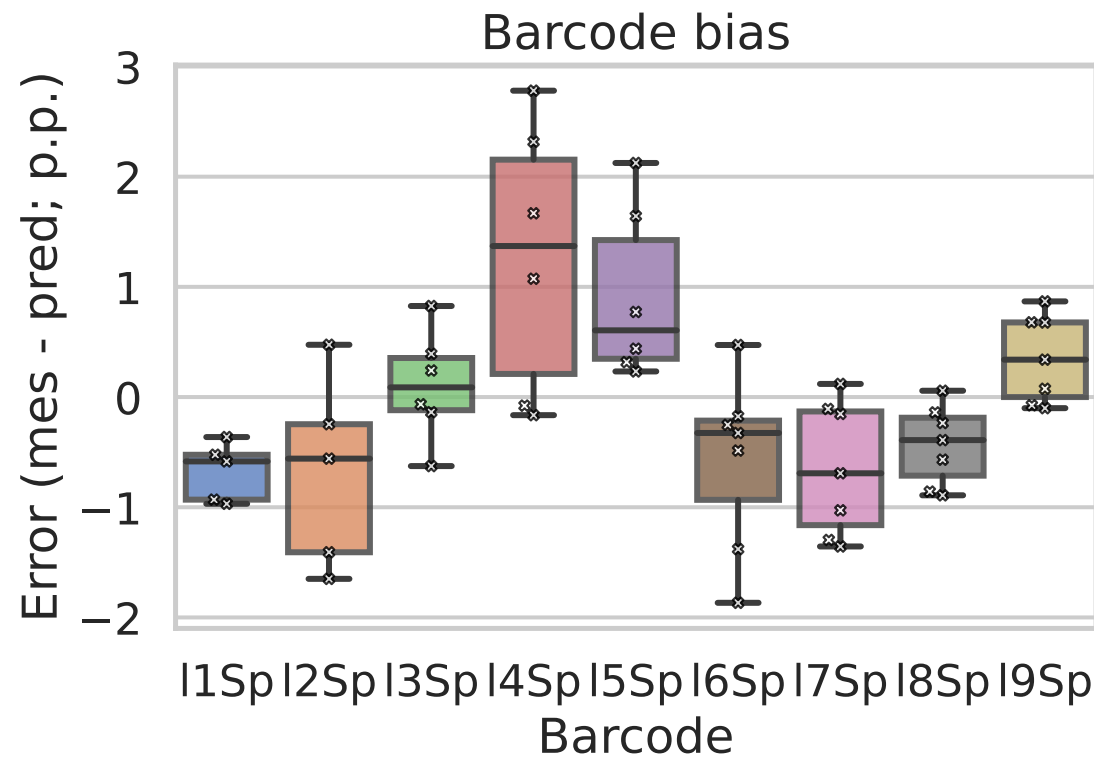
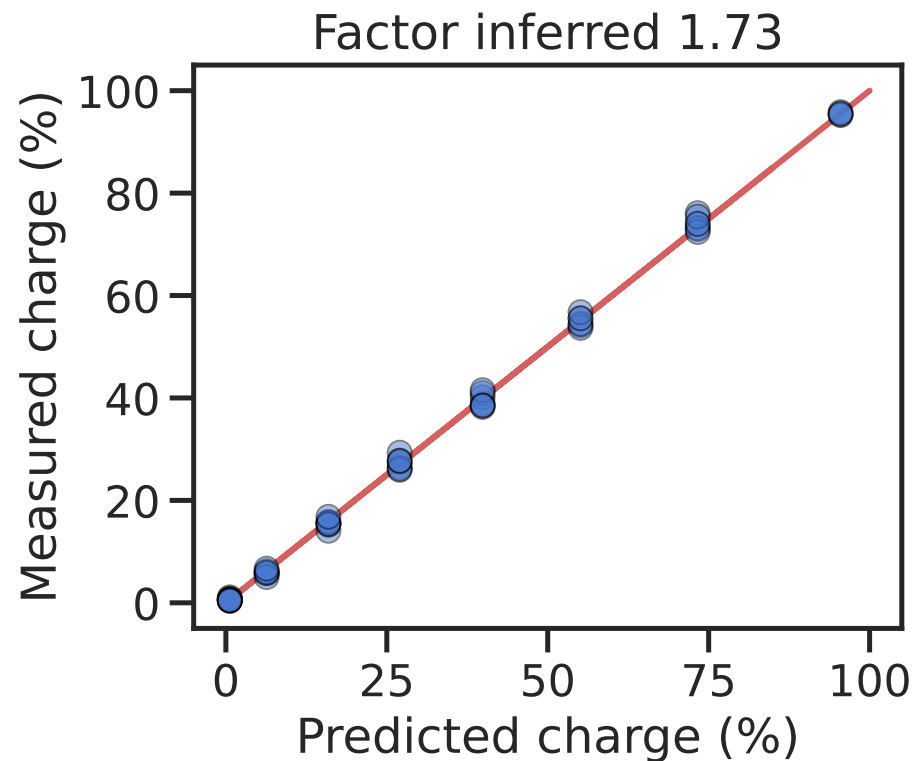
Ser-GCT-1-1



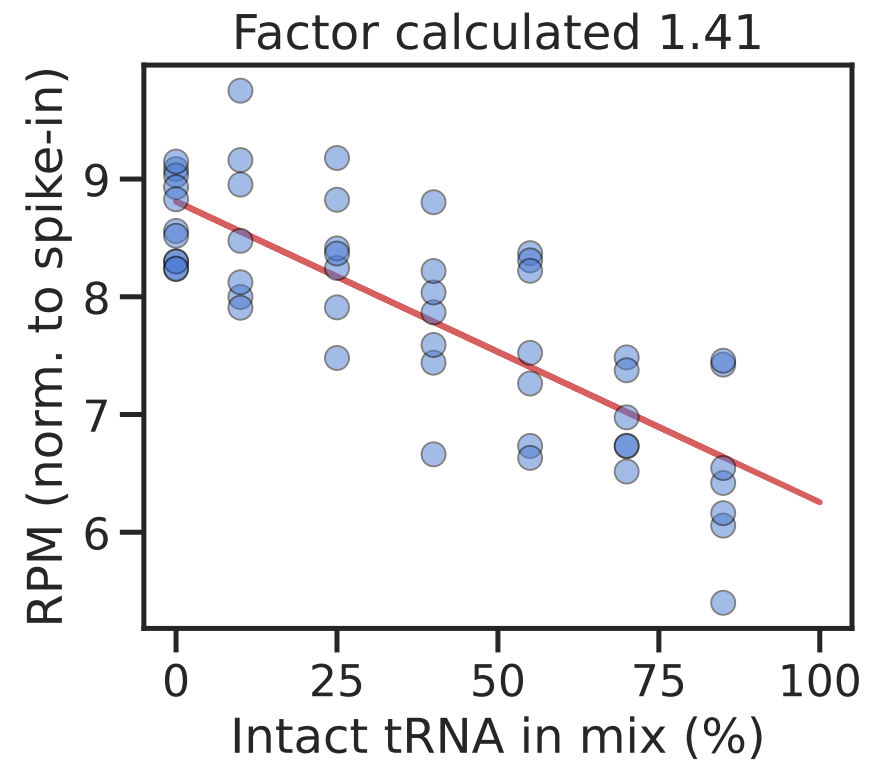
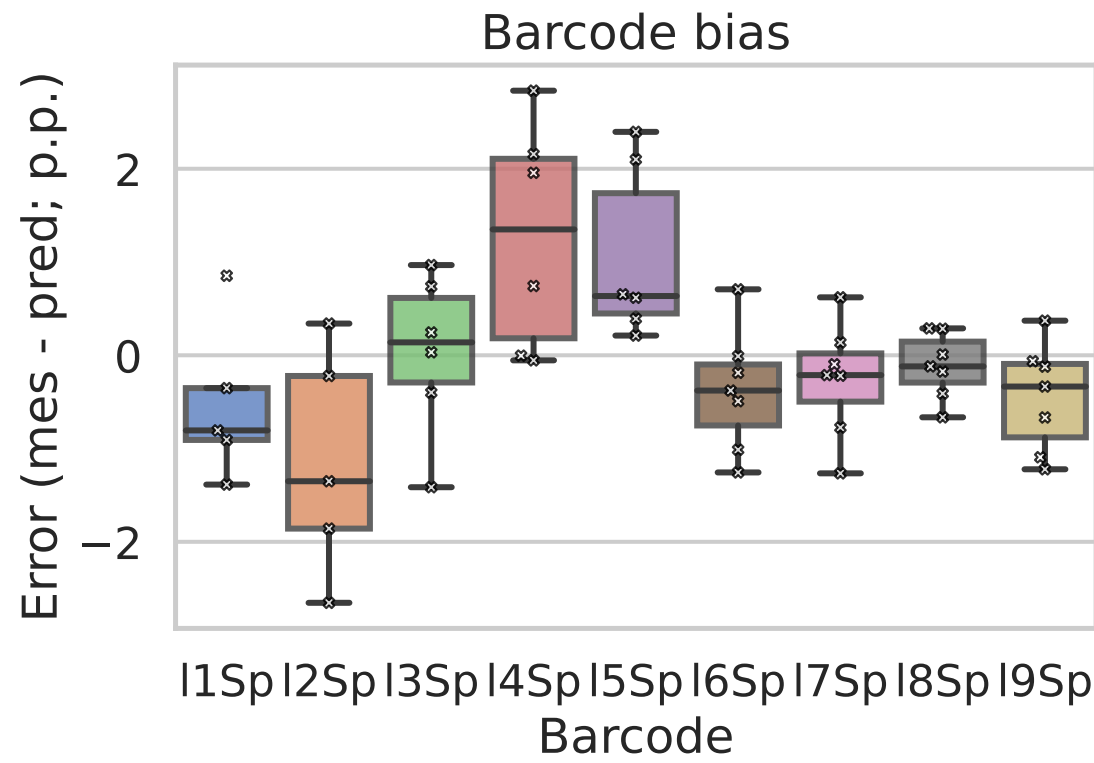
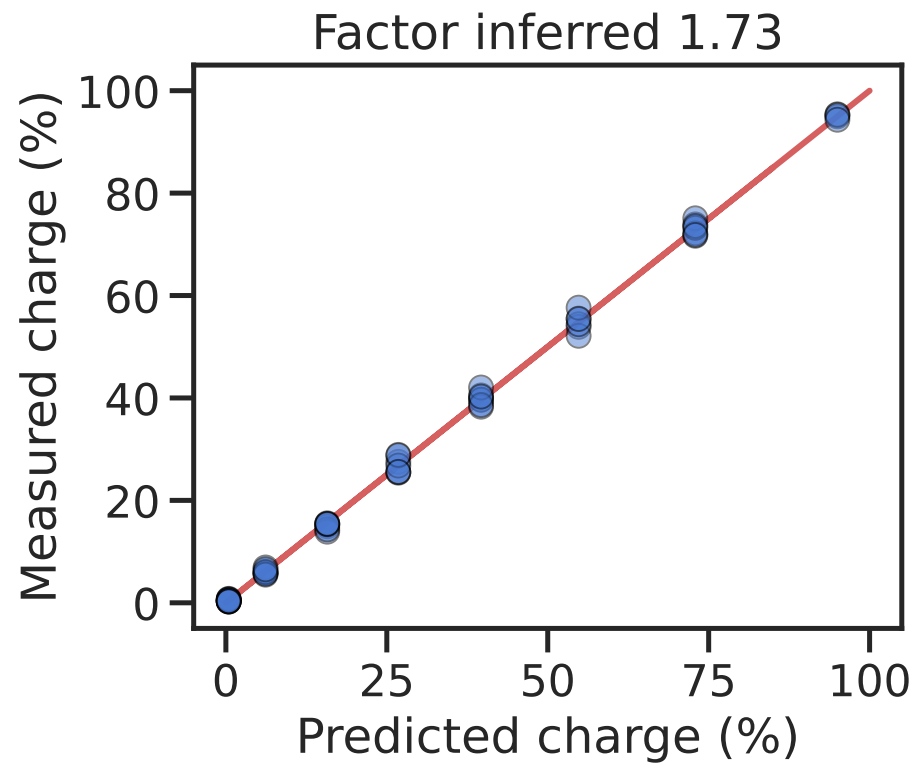
Met-CAT-1-1



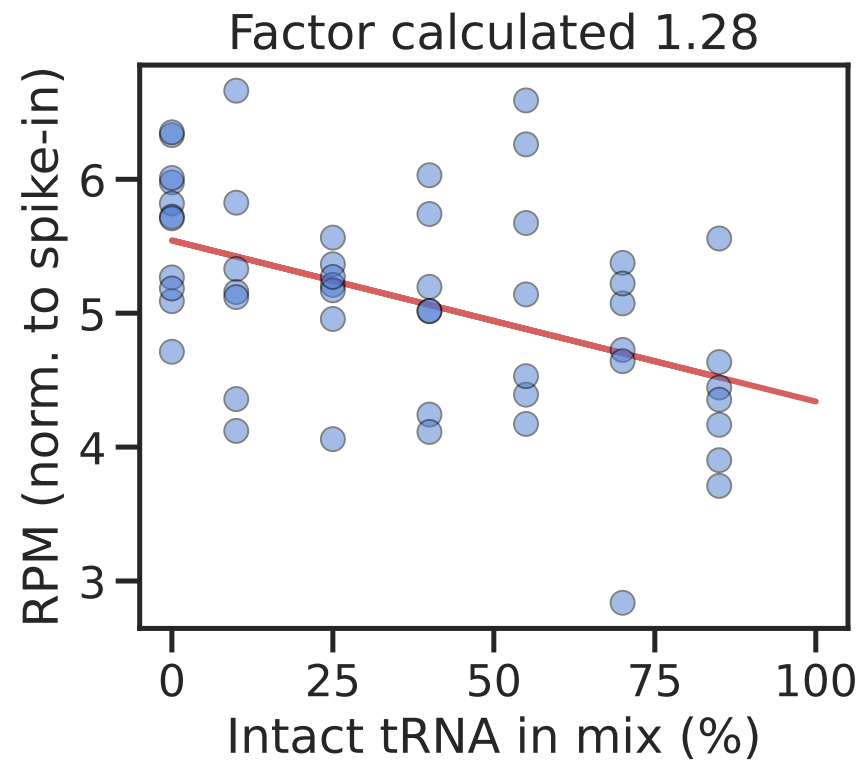
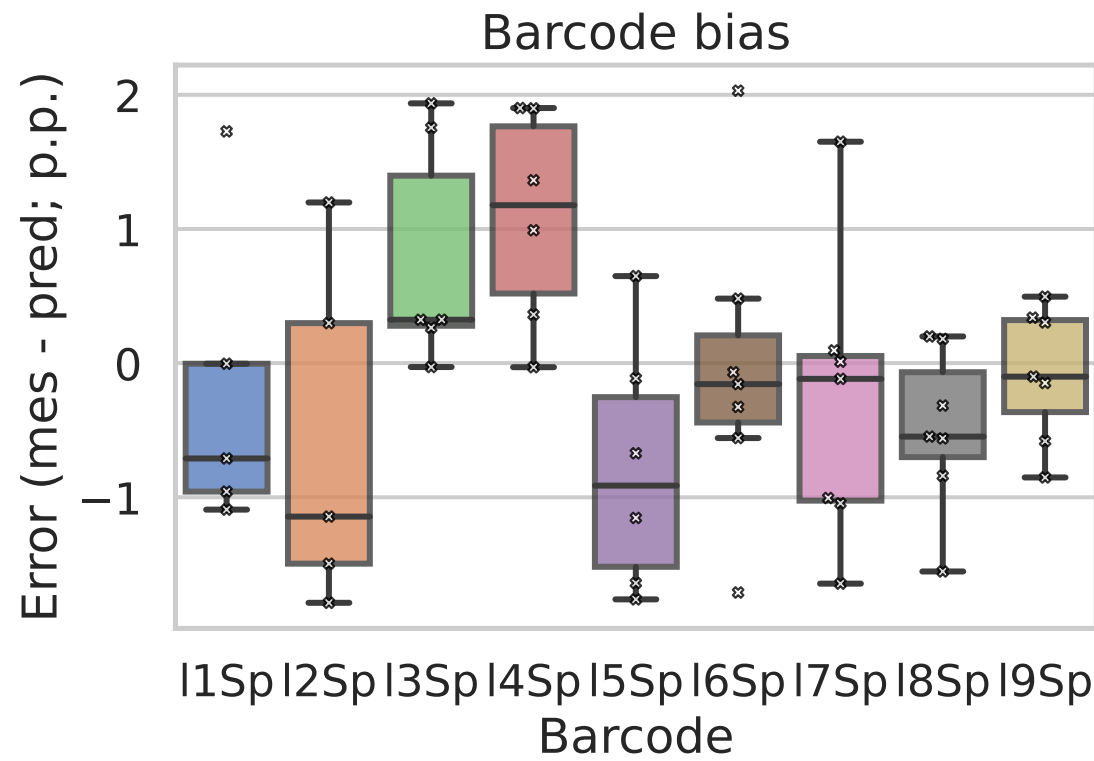
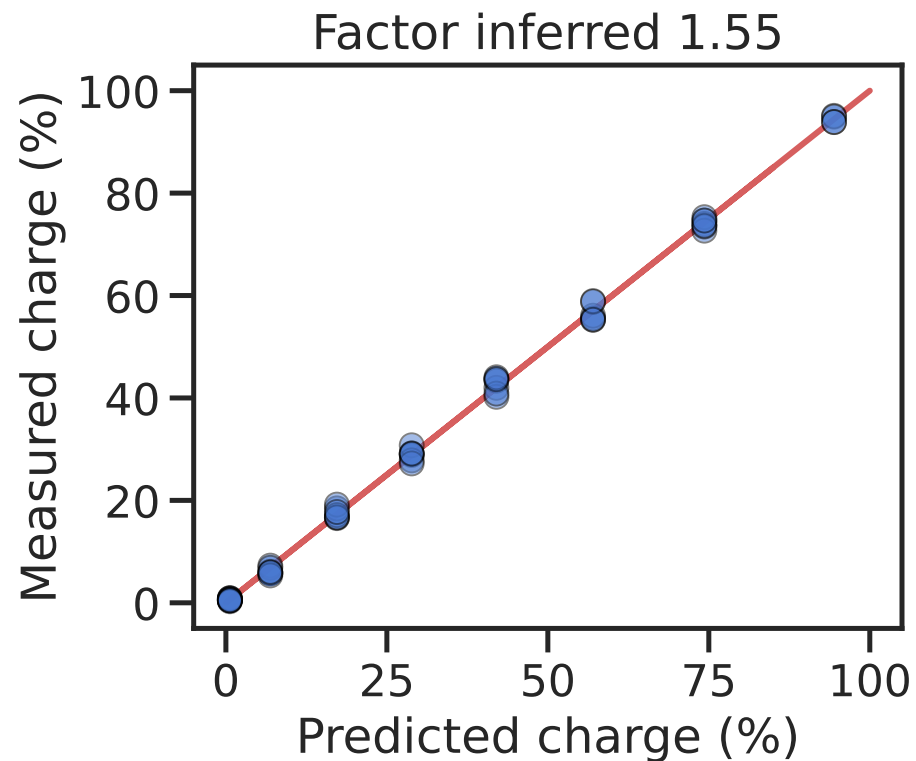
Arg-TCT-2-1



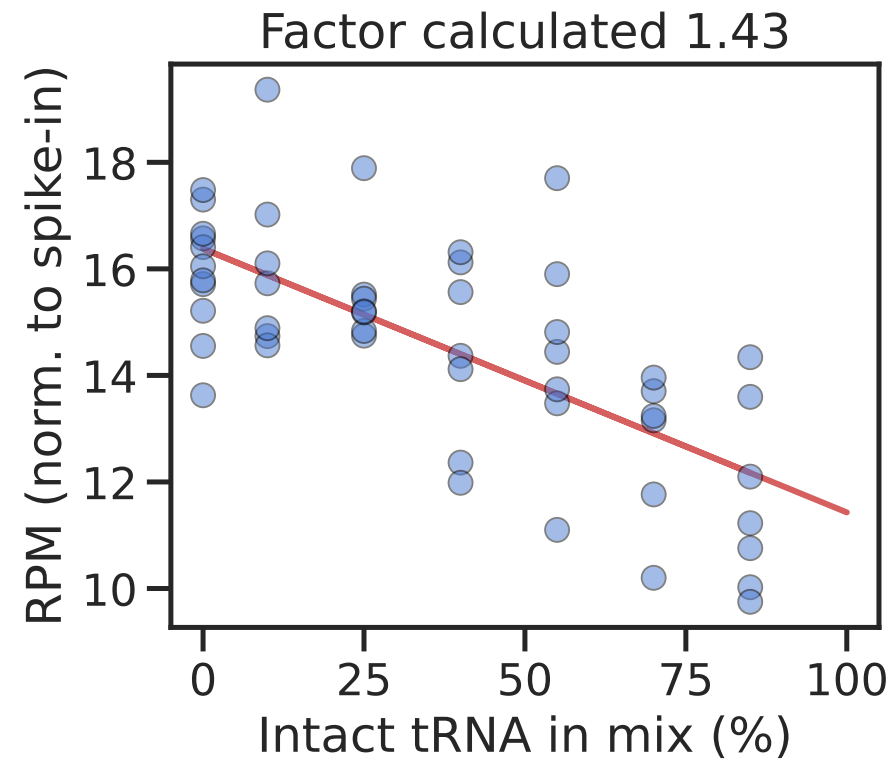
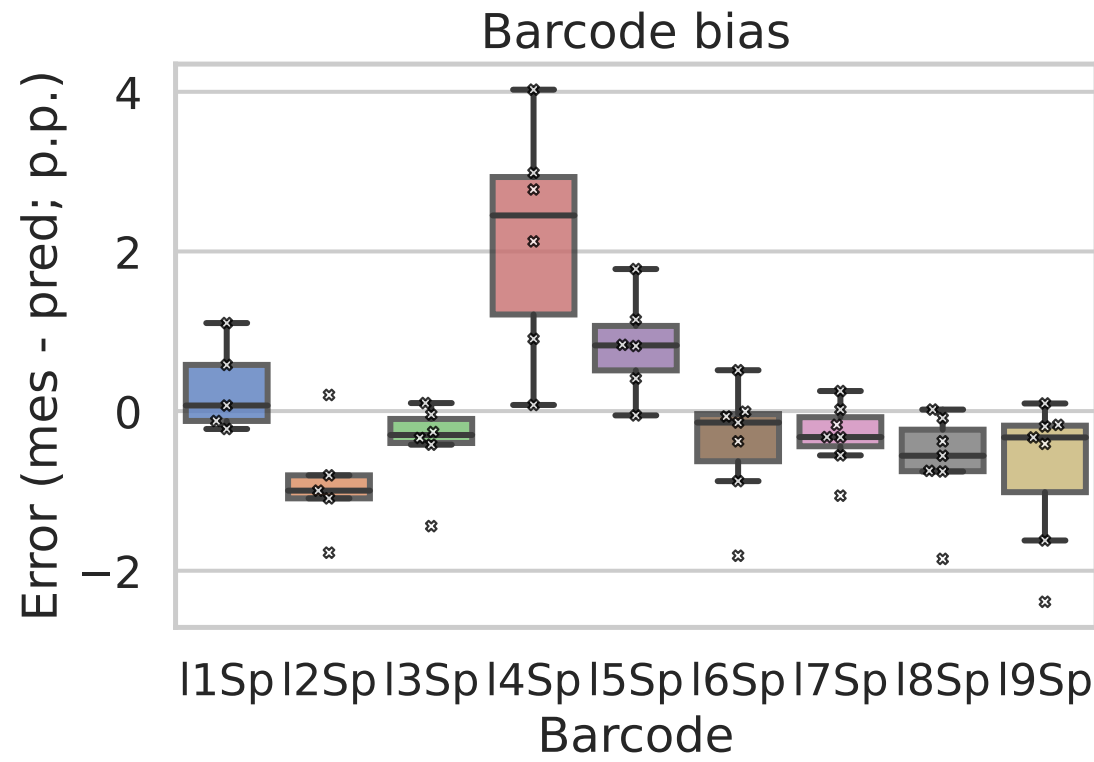
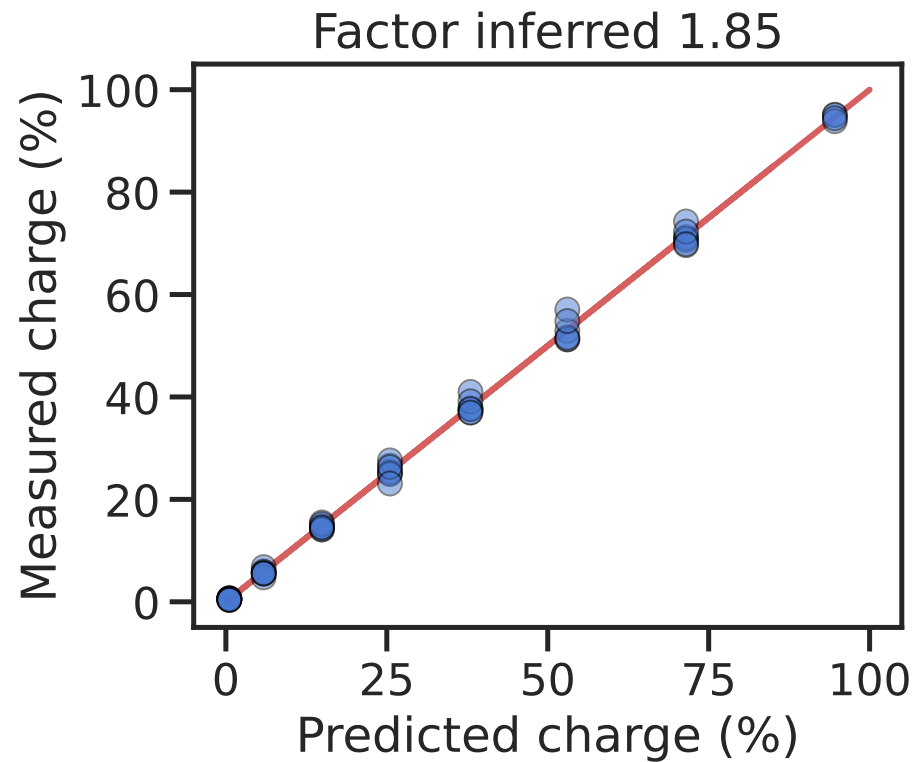
Arg-TCT-4-1



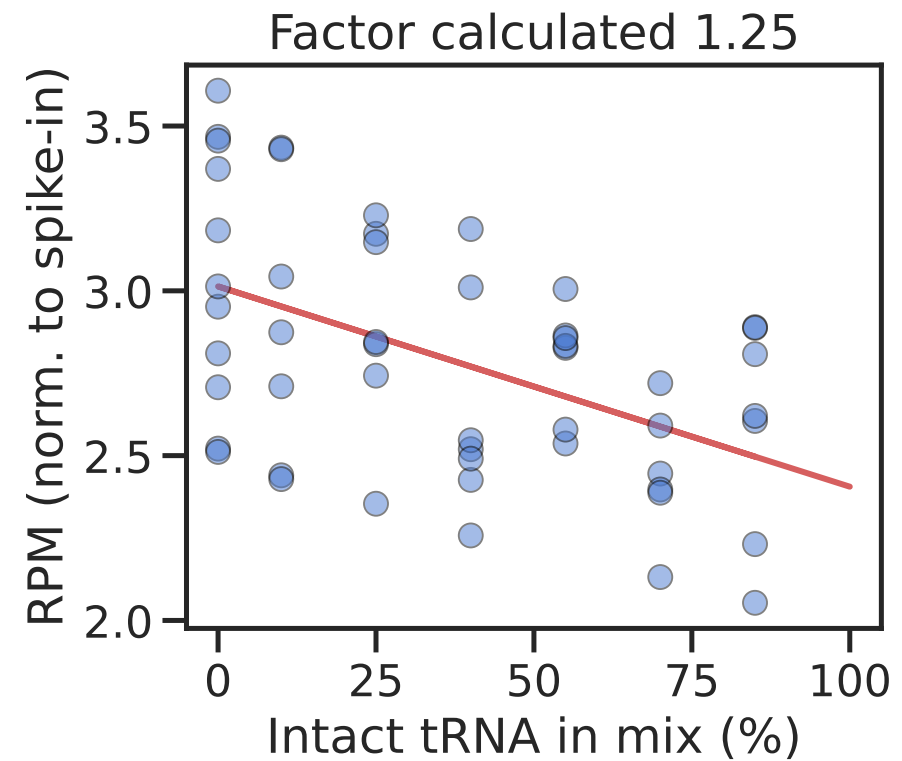
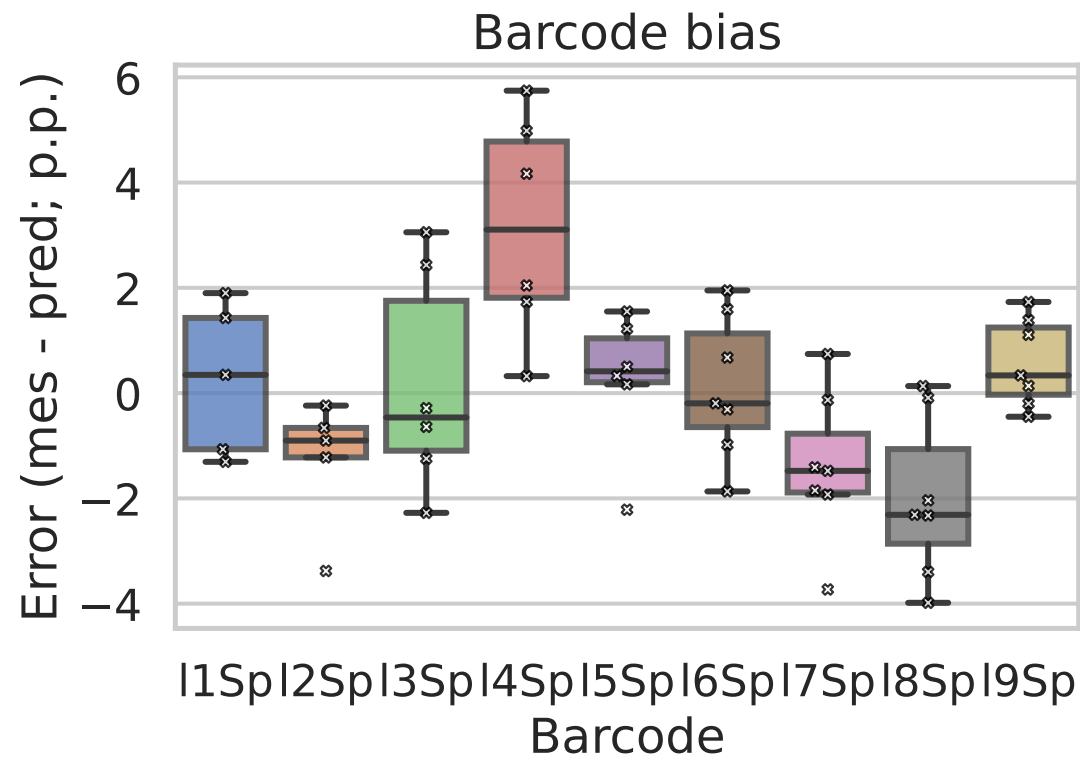
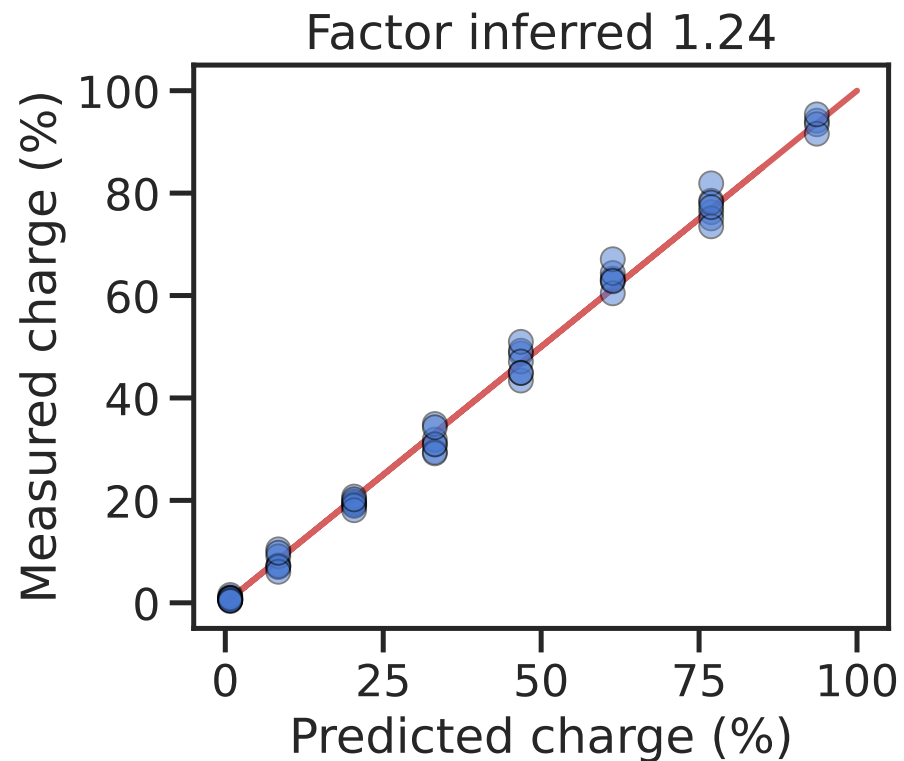
Leu-AAG-1-1



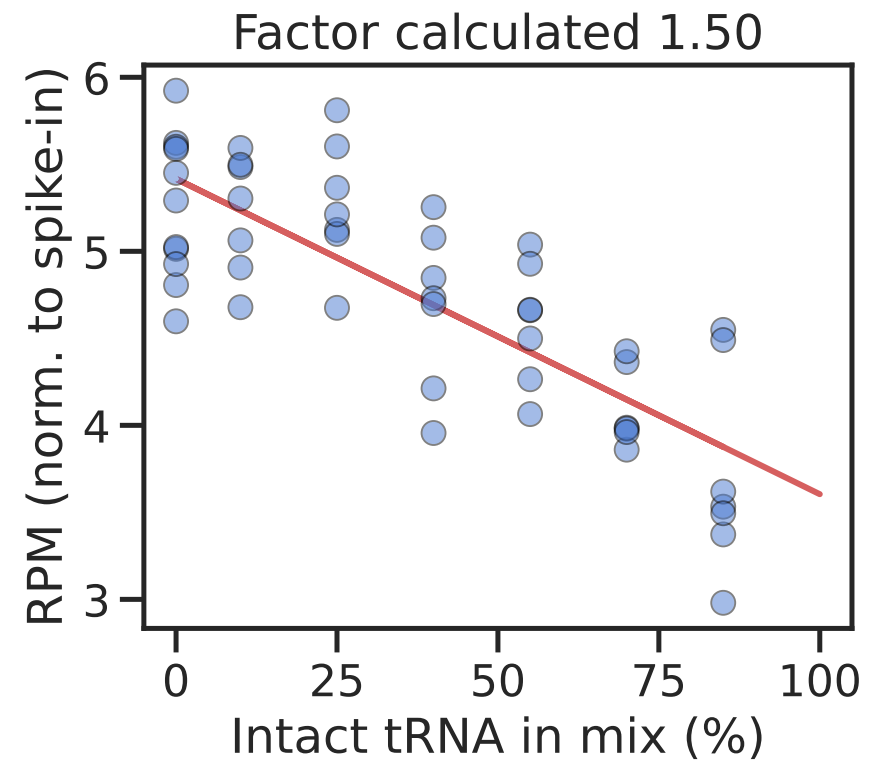
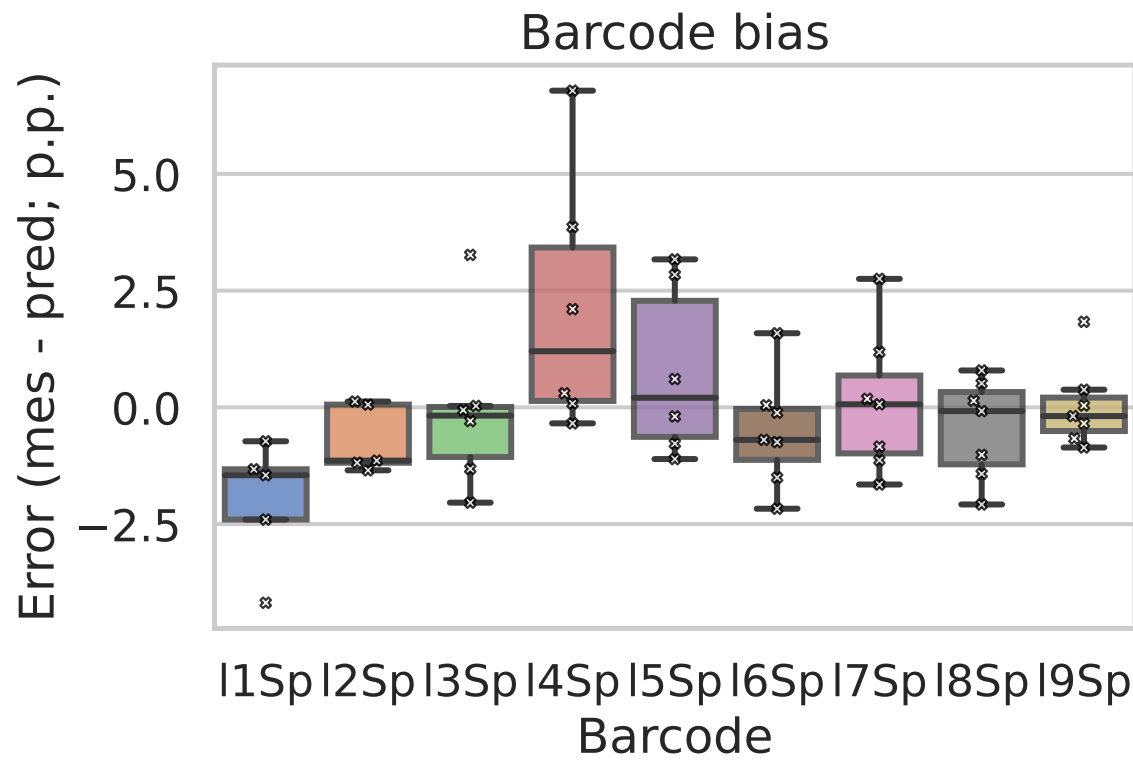
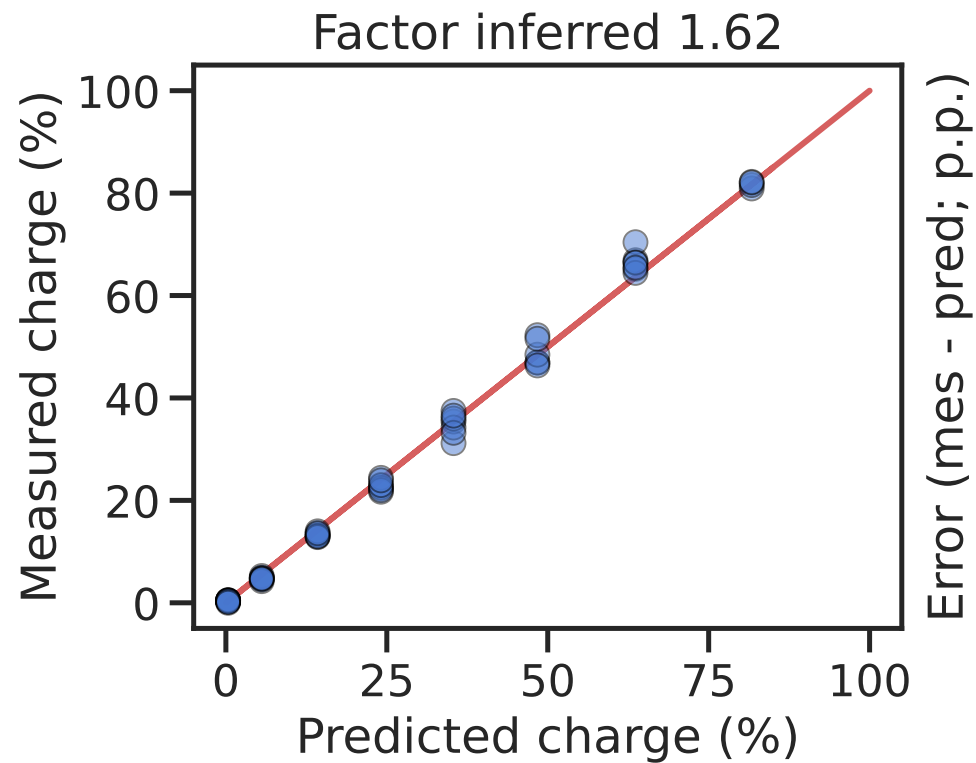
Leu-CAG-2-1



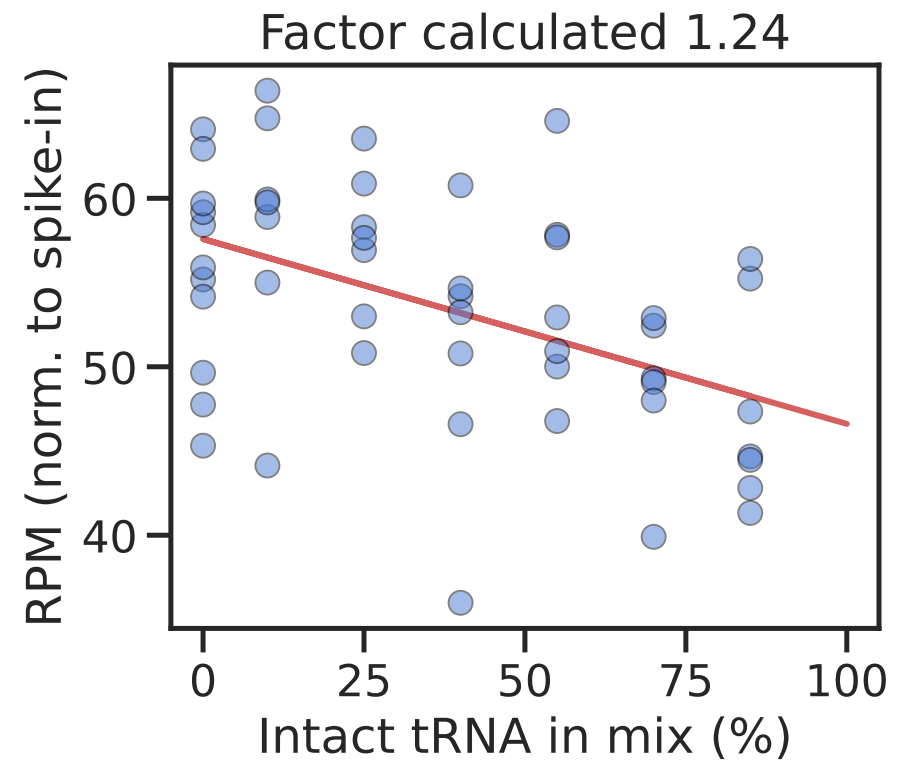
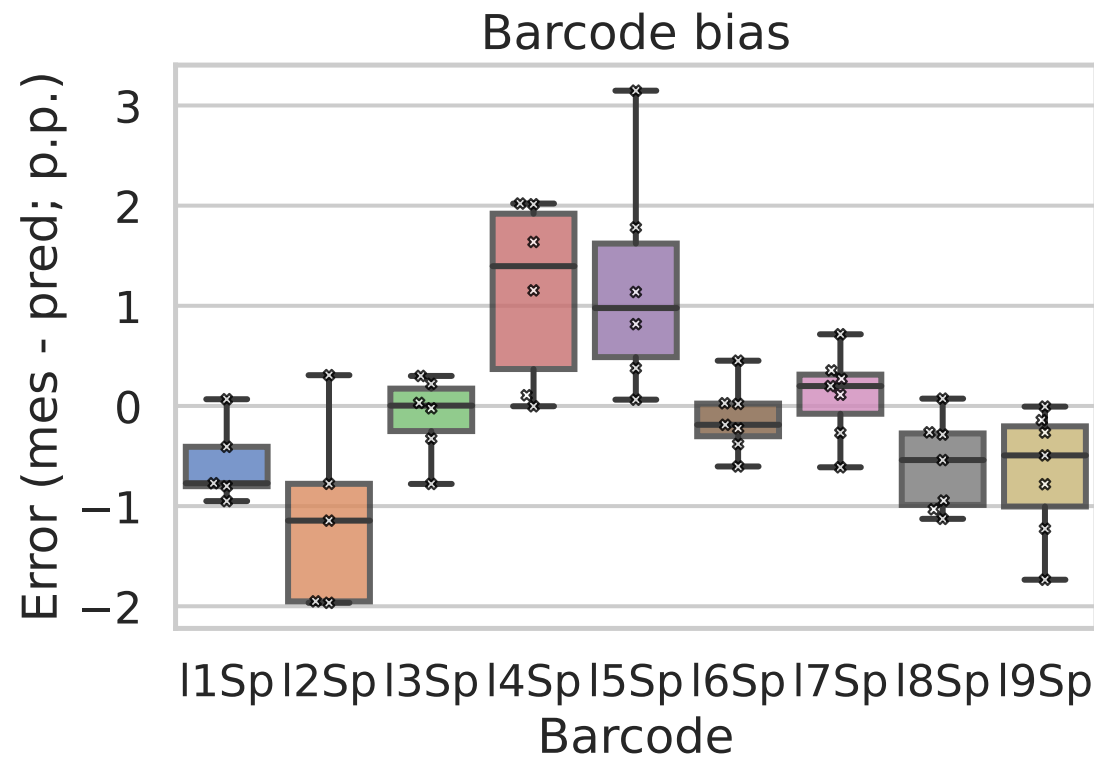
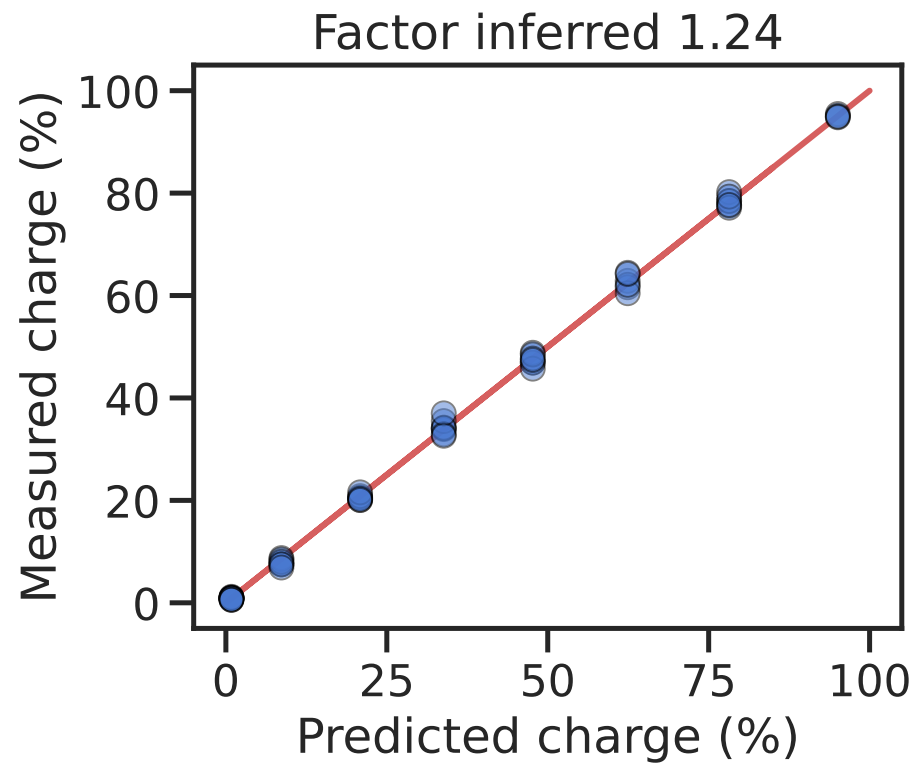
Thr-AGT-6-1

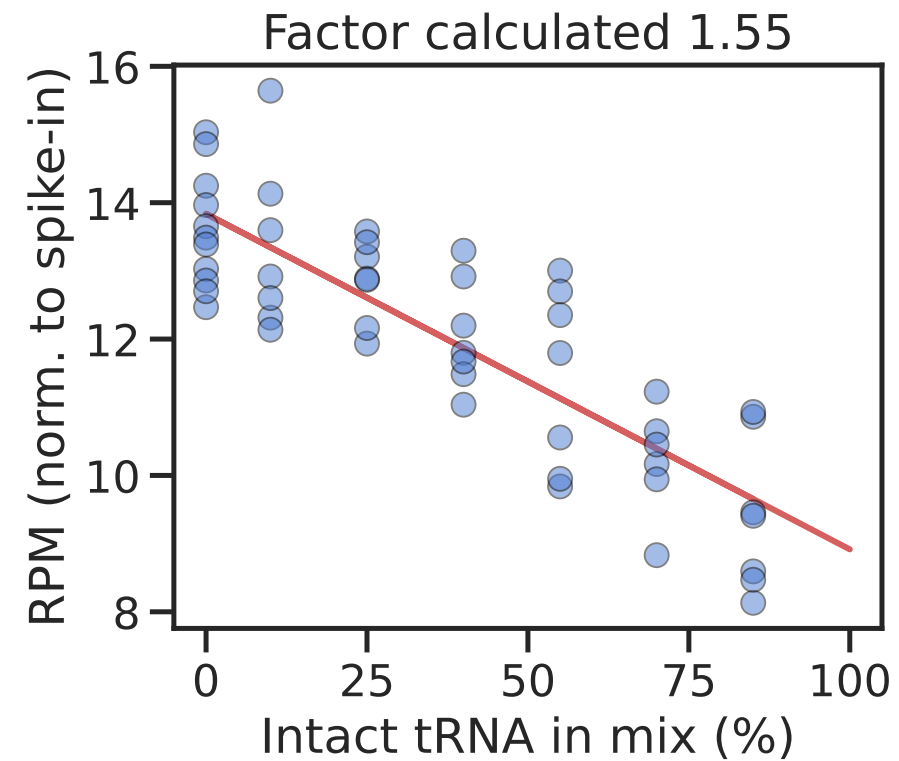
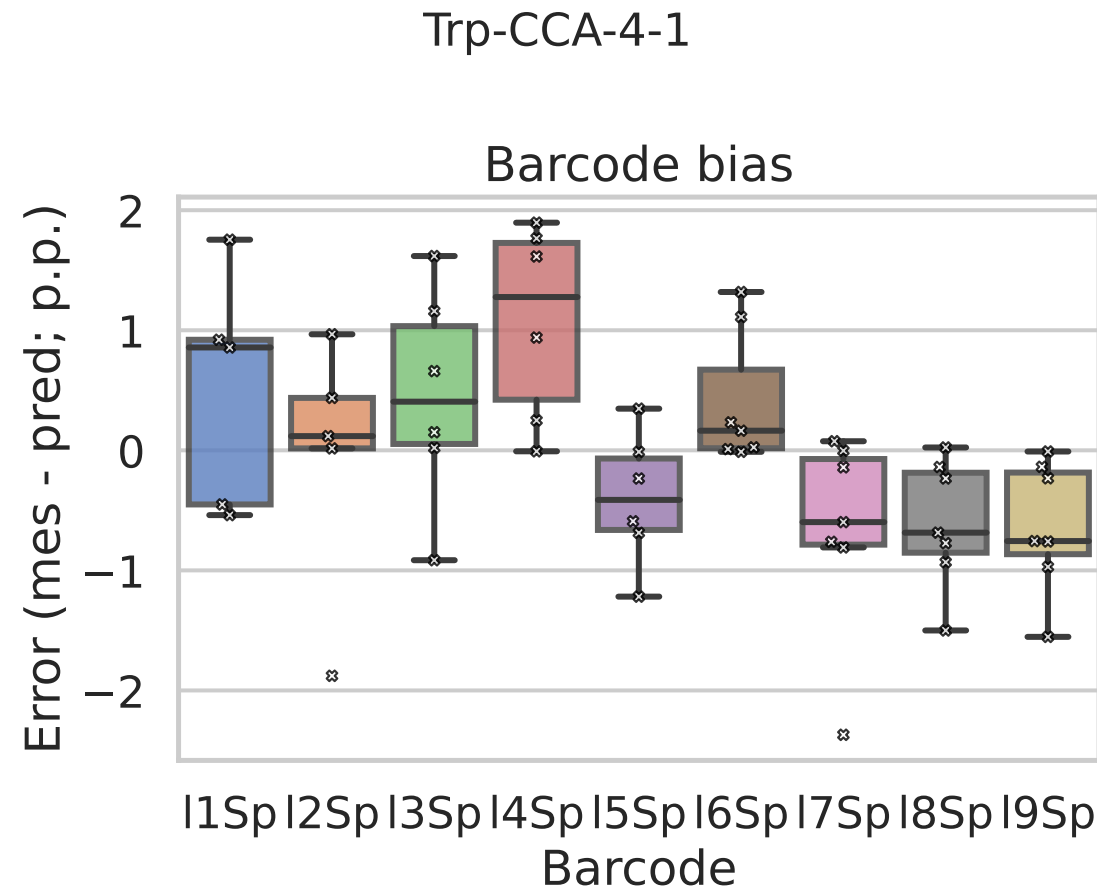
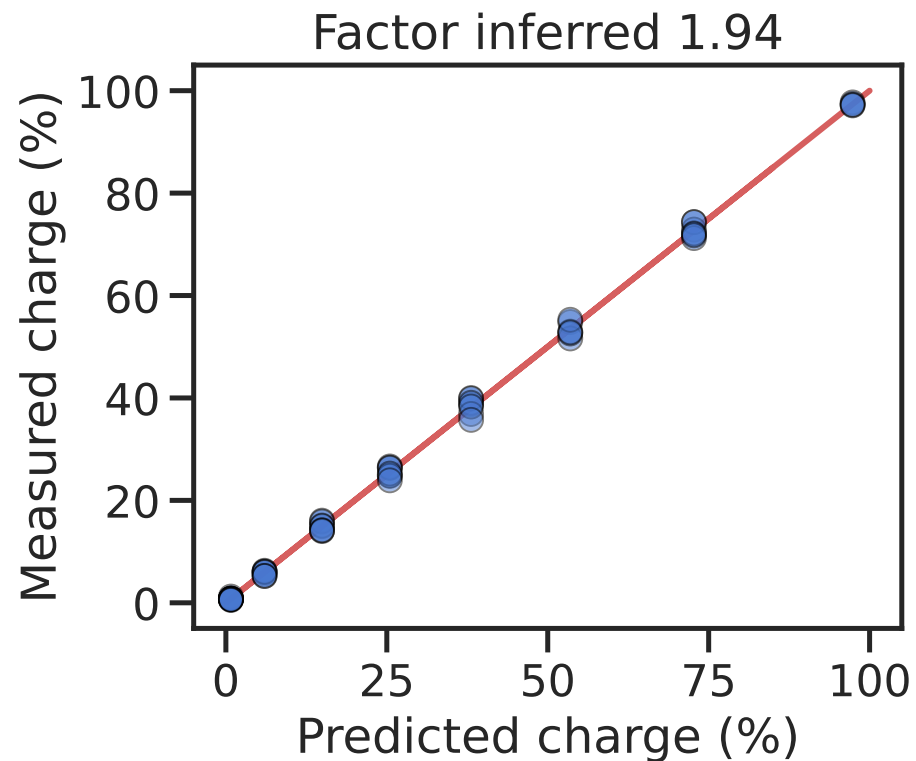


Ser-GCT-5-1

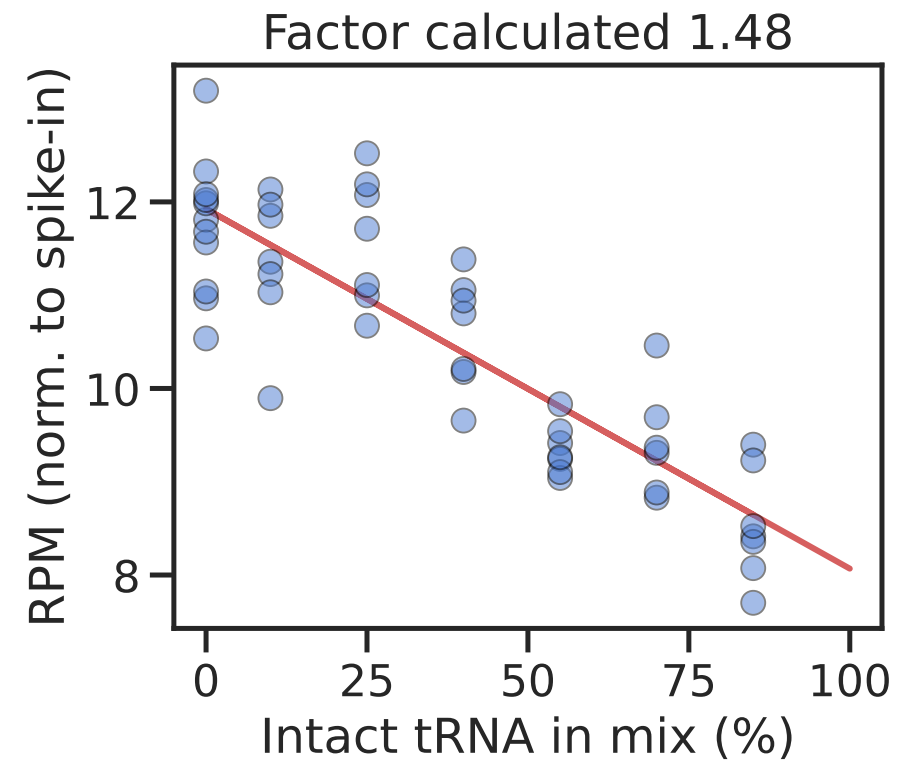
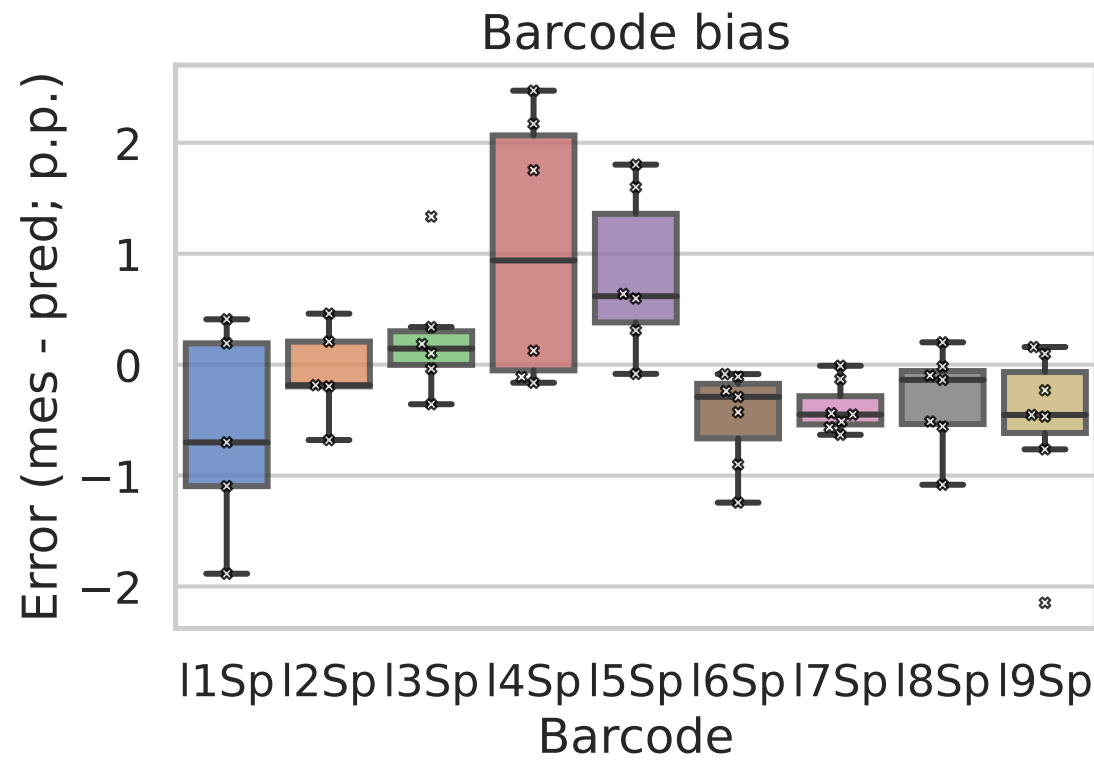
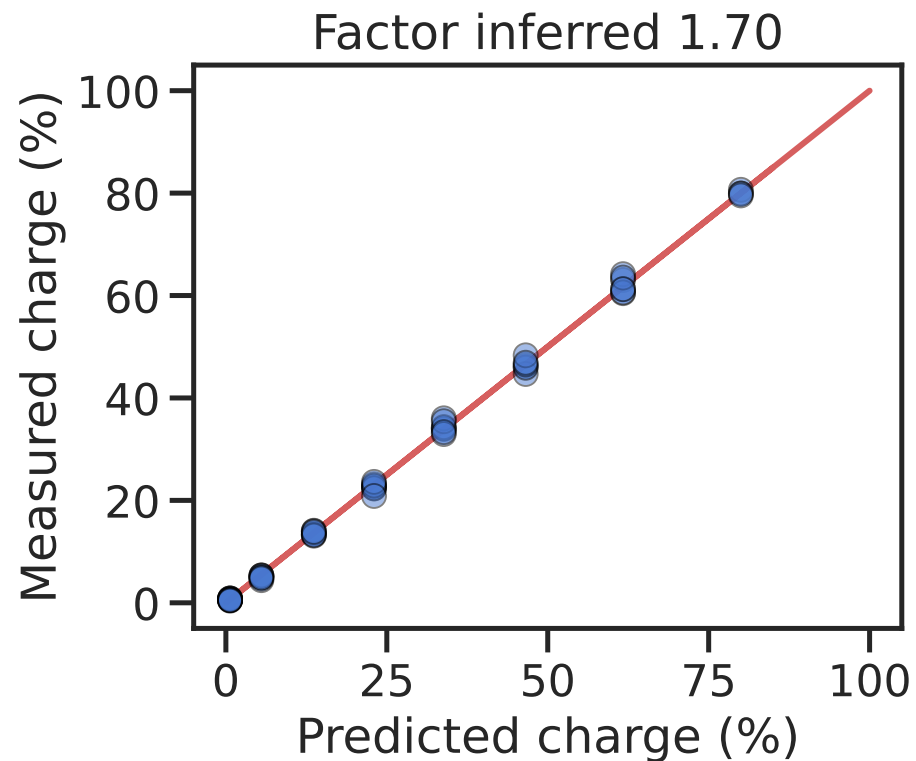


Asp-GTC-2-1

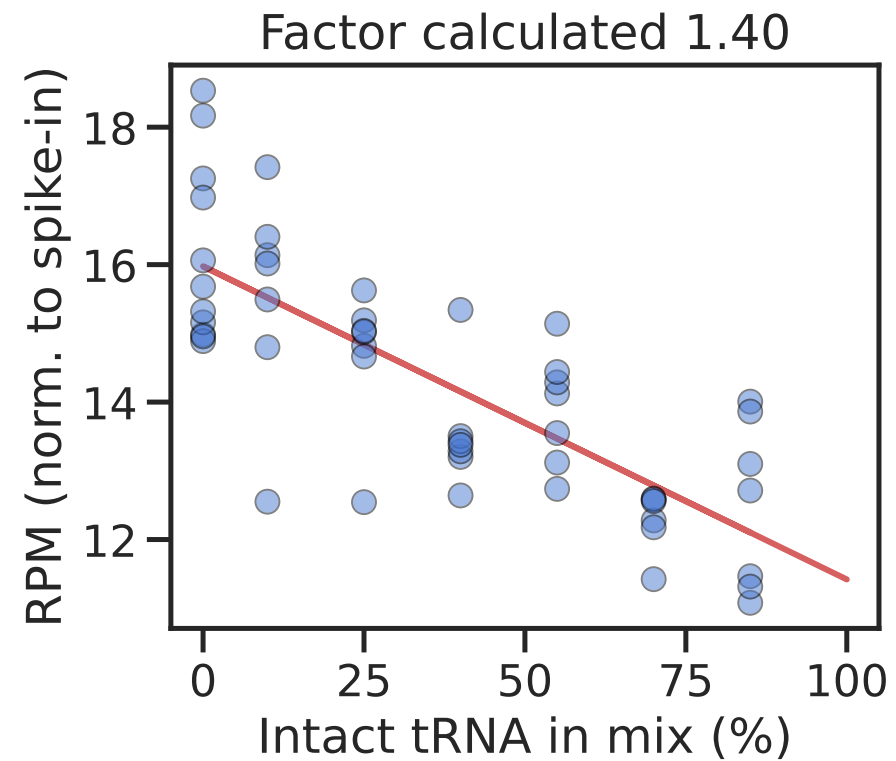
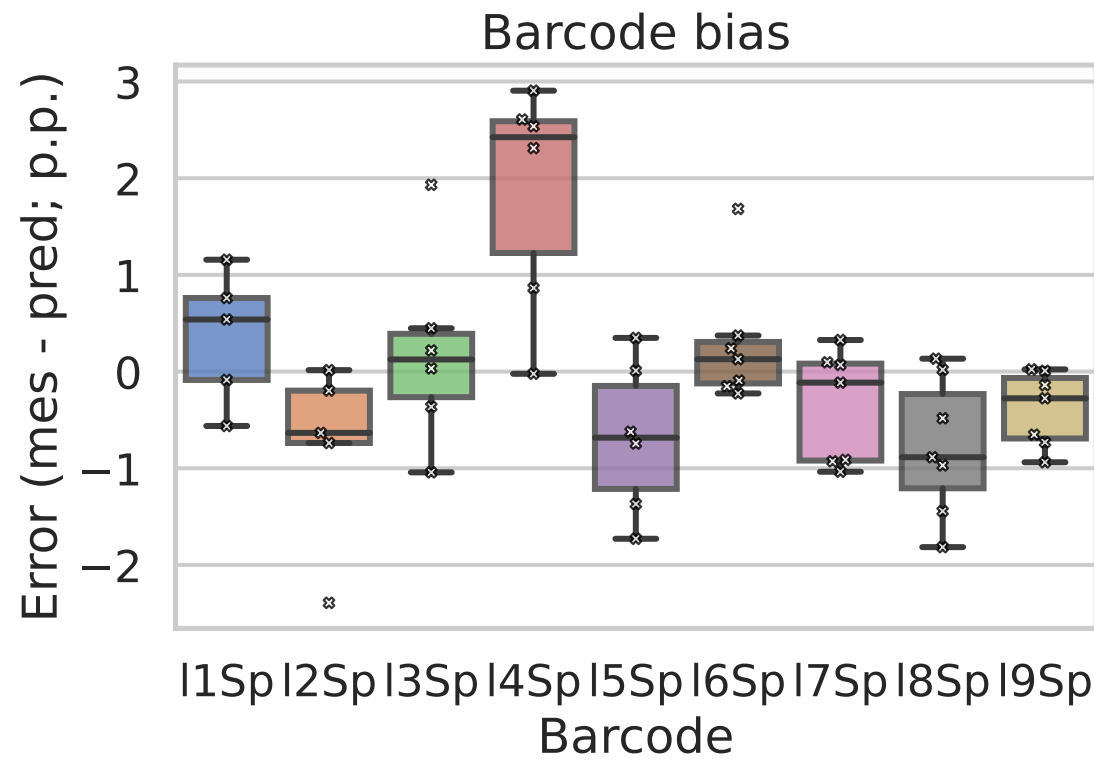
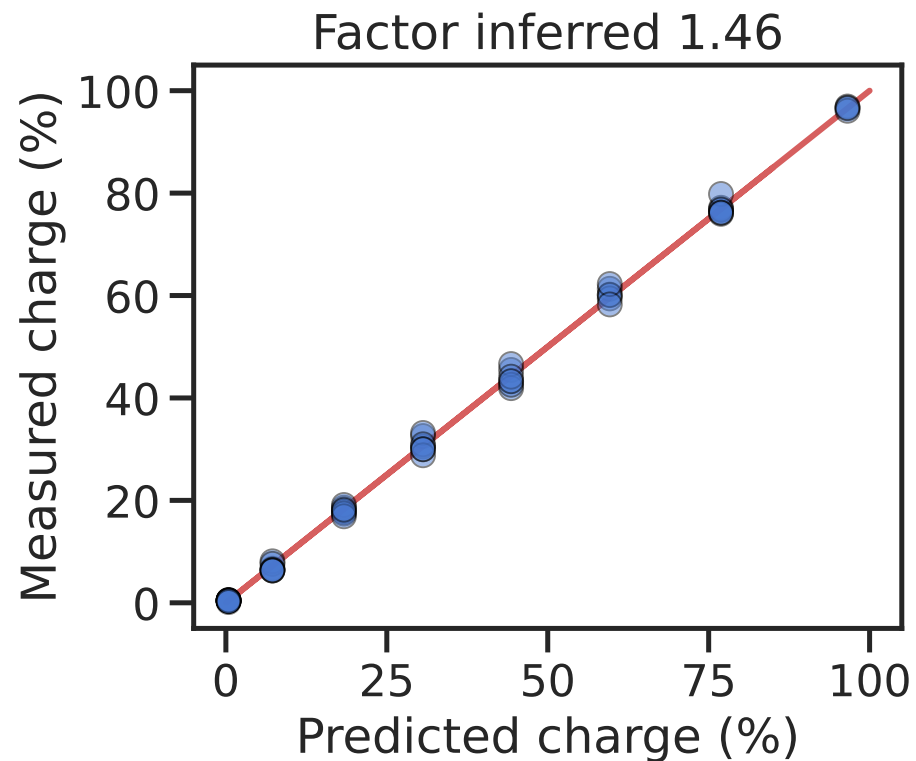




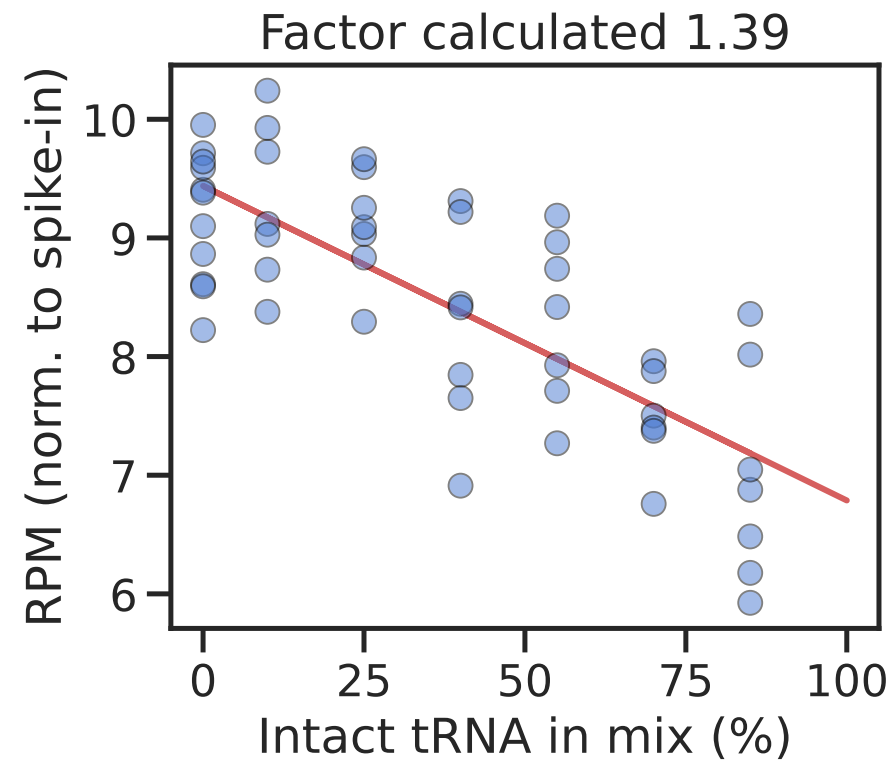
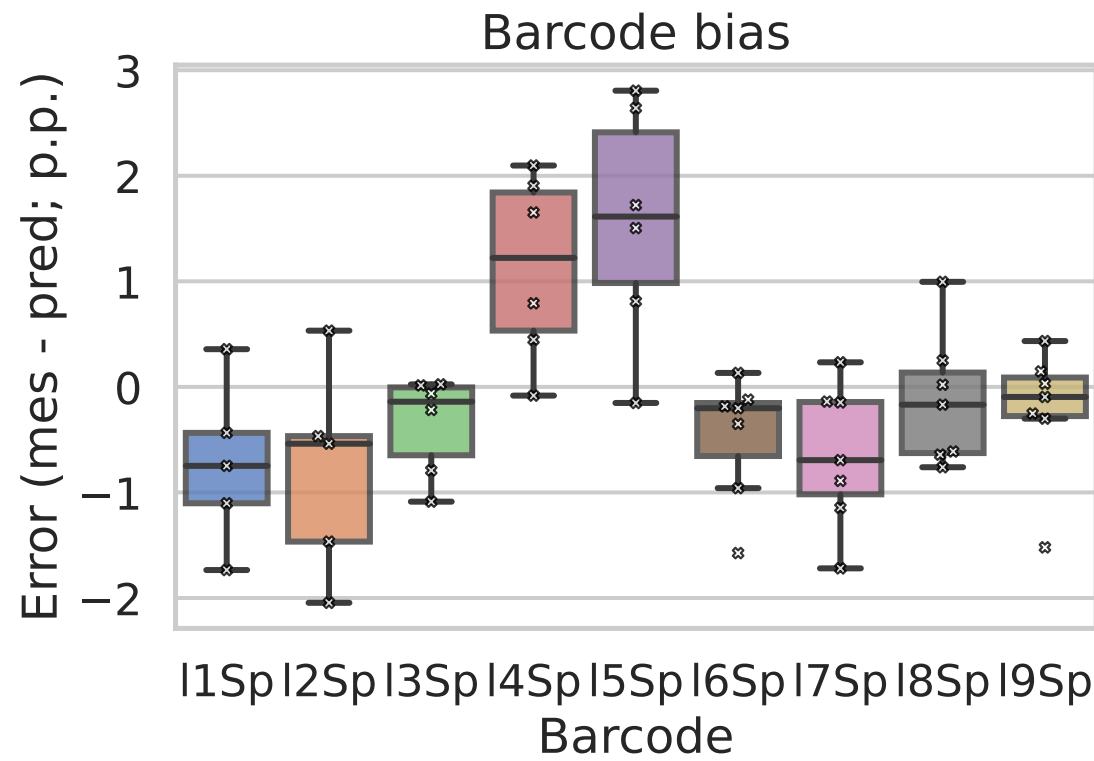
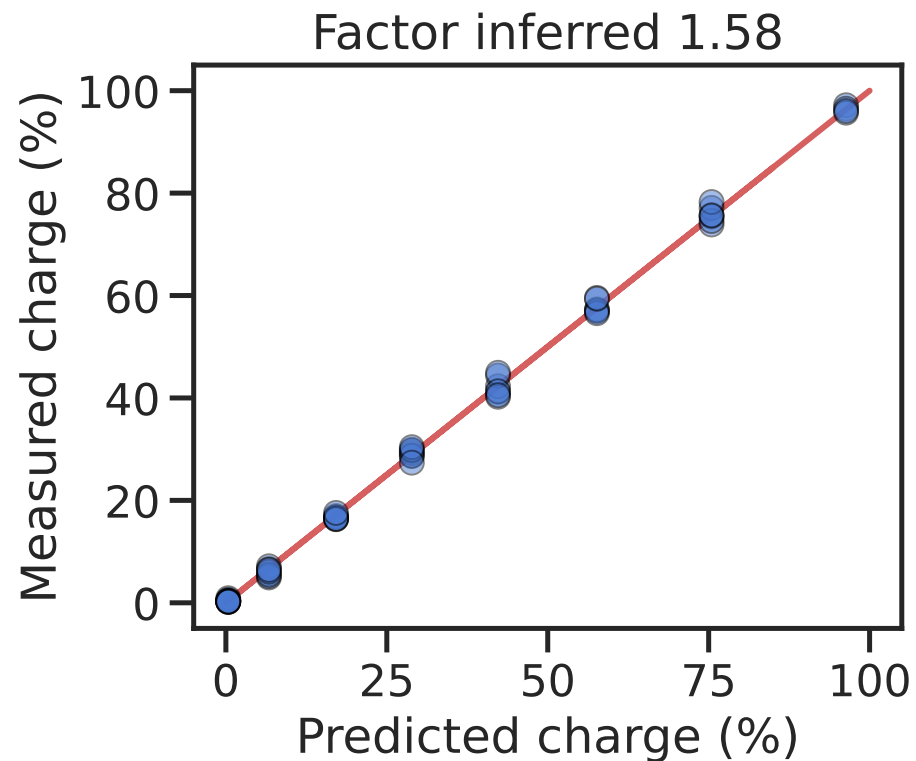
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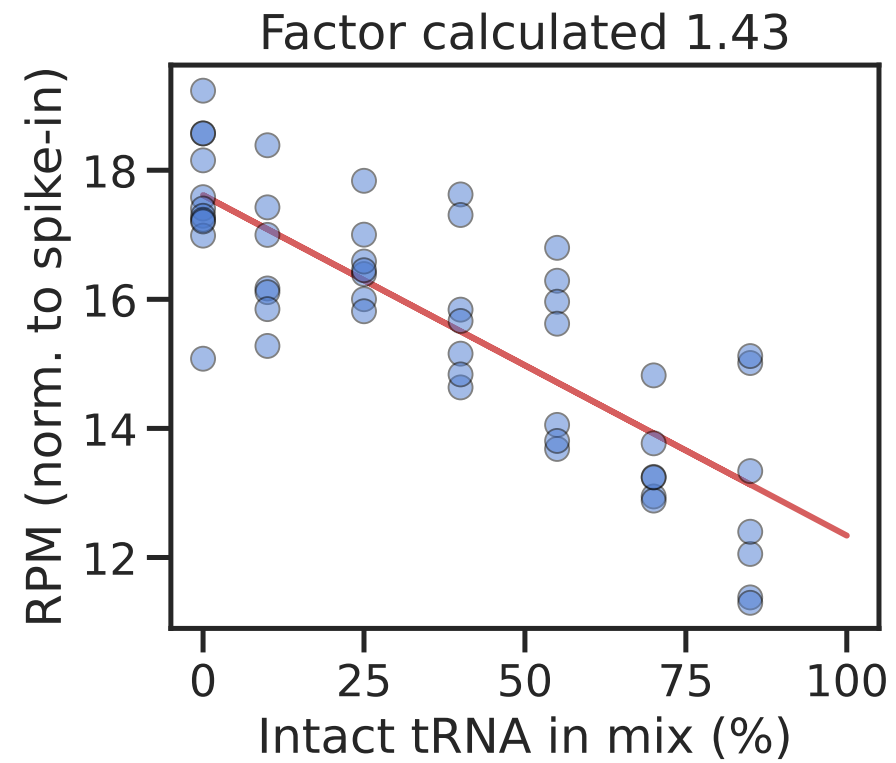
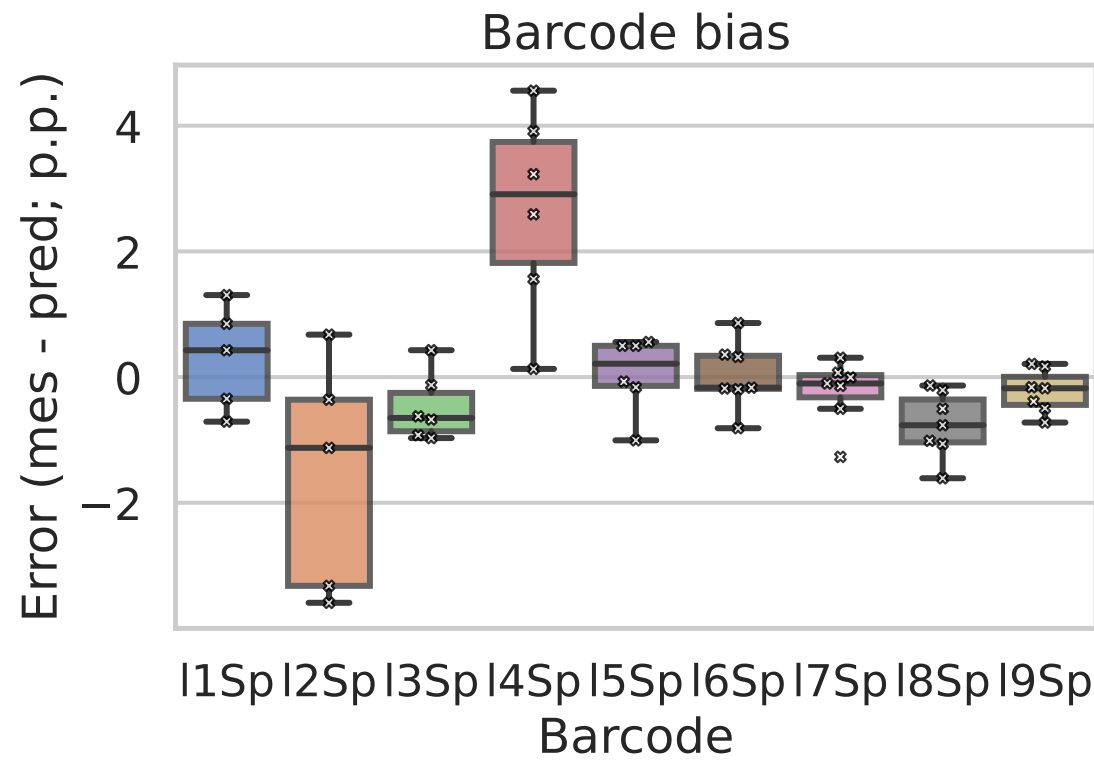
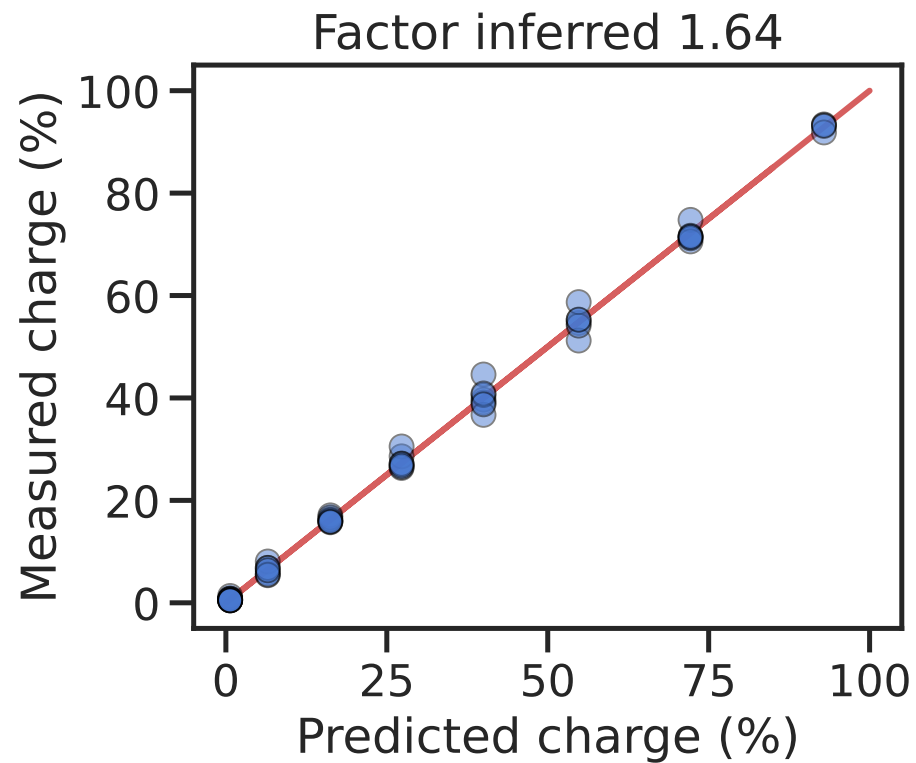
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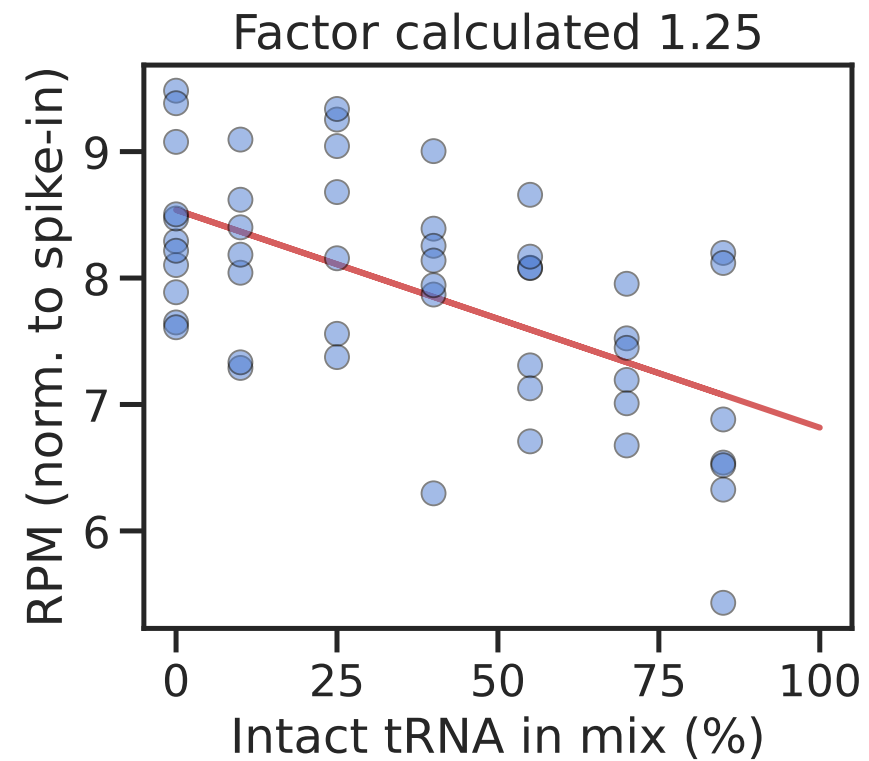
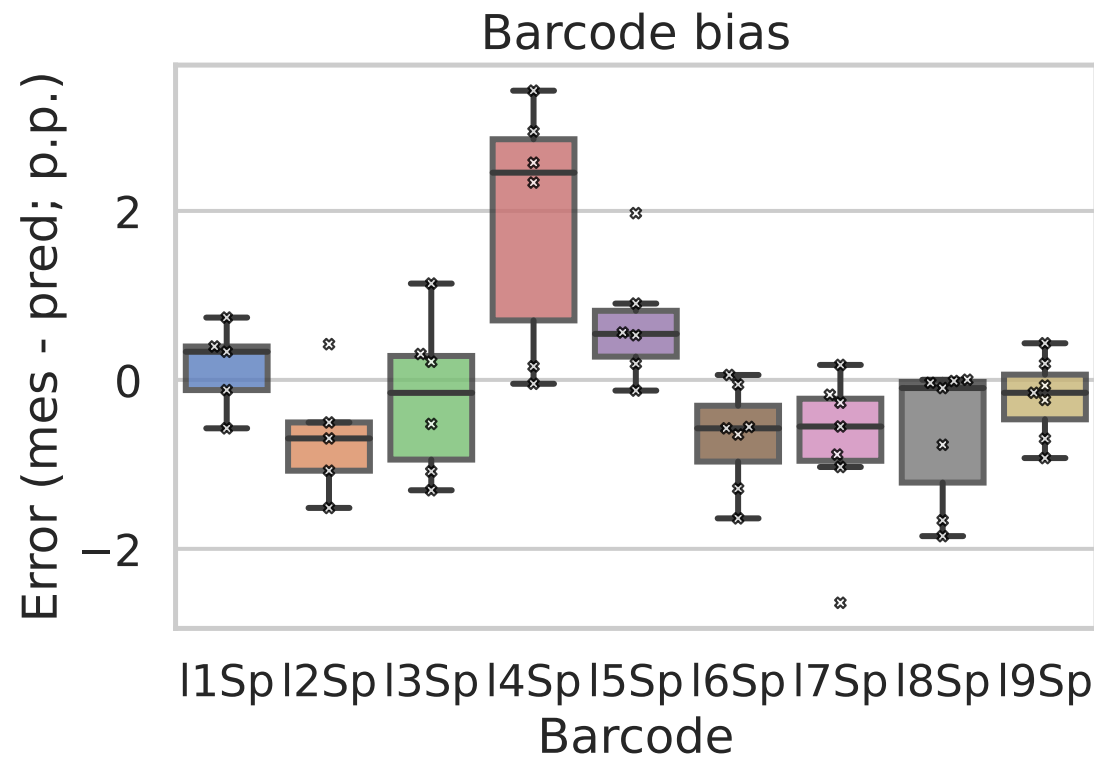
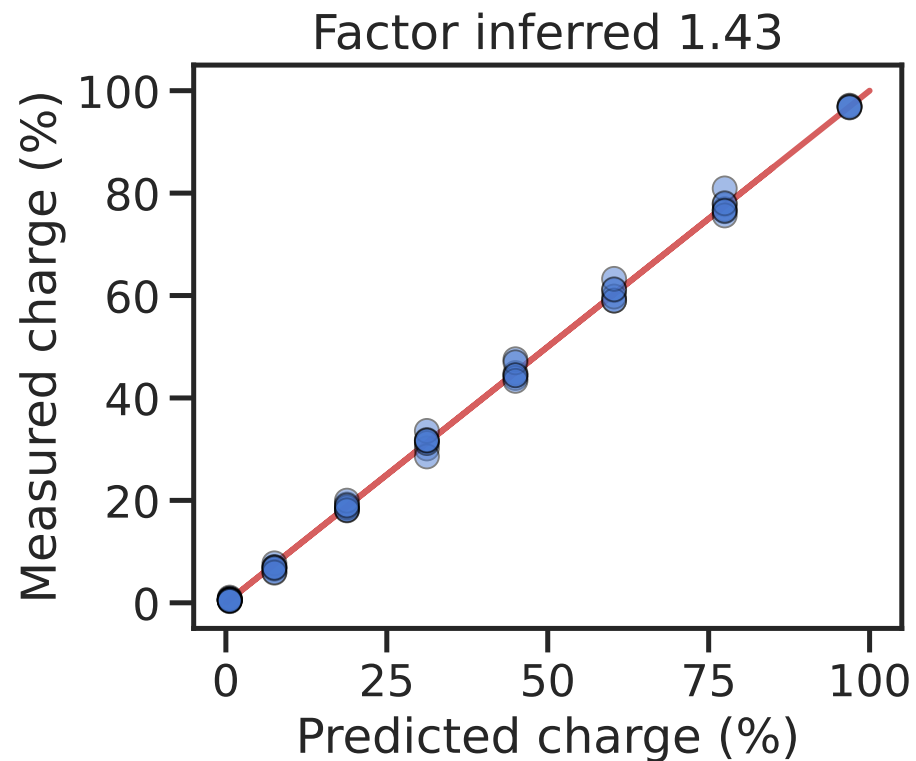
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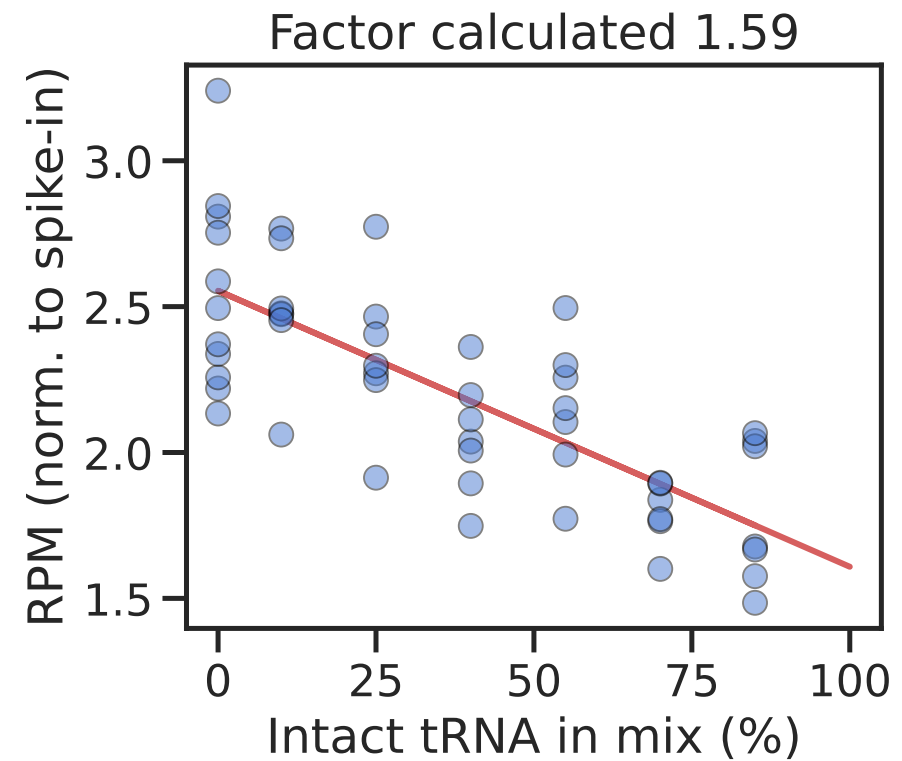
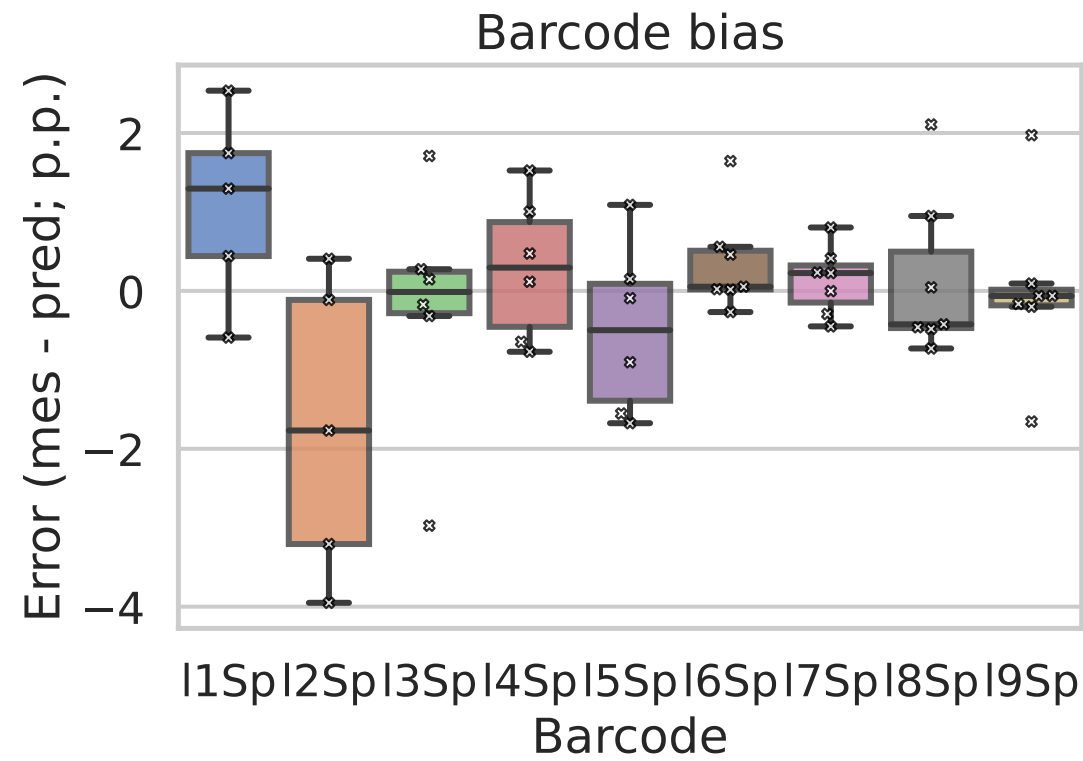
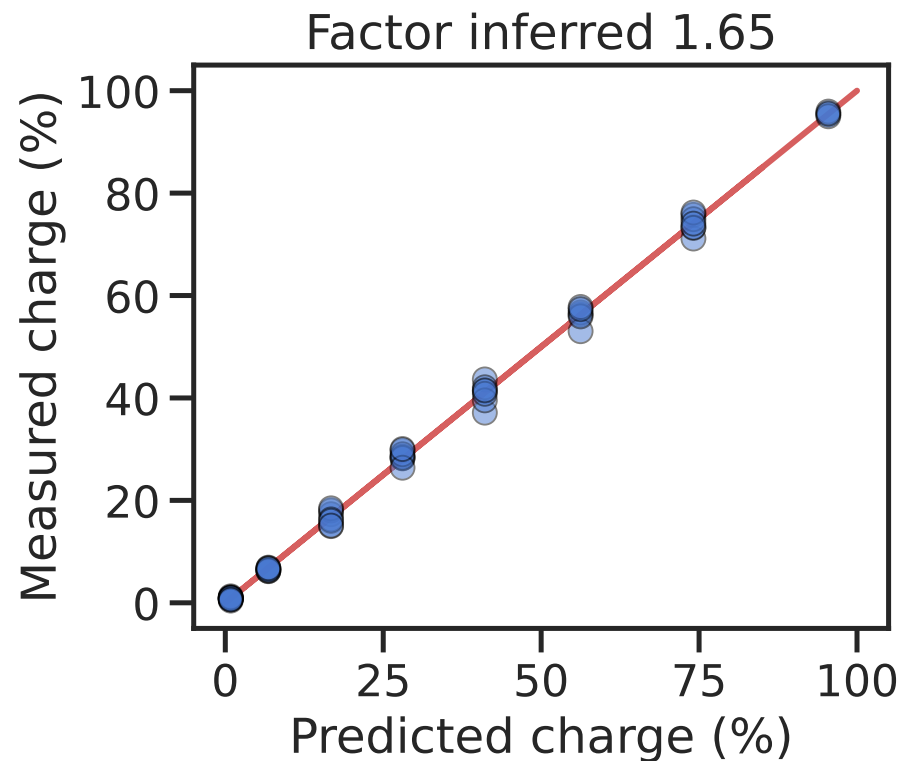
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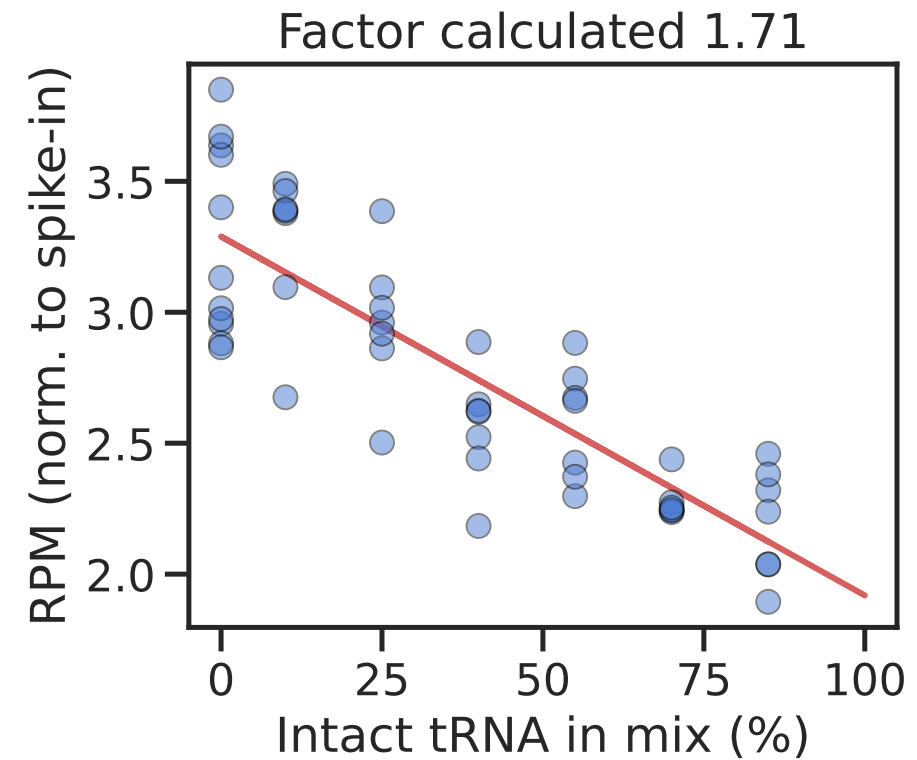
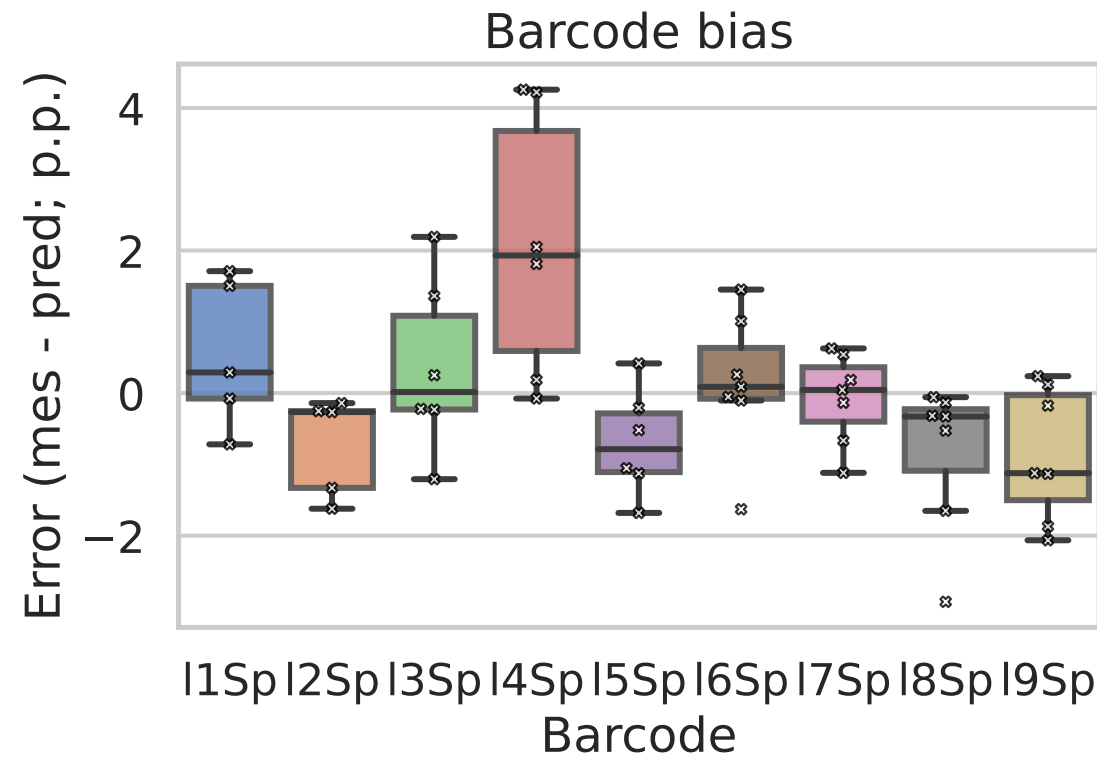
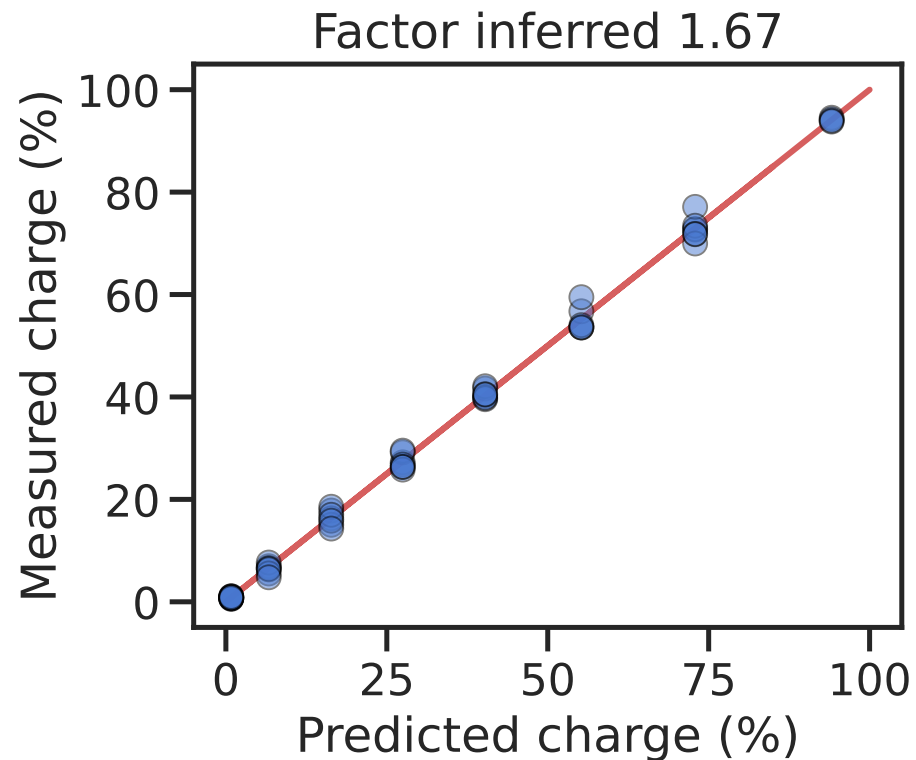
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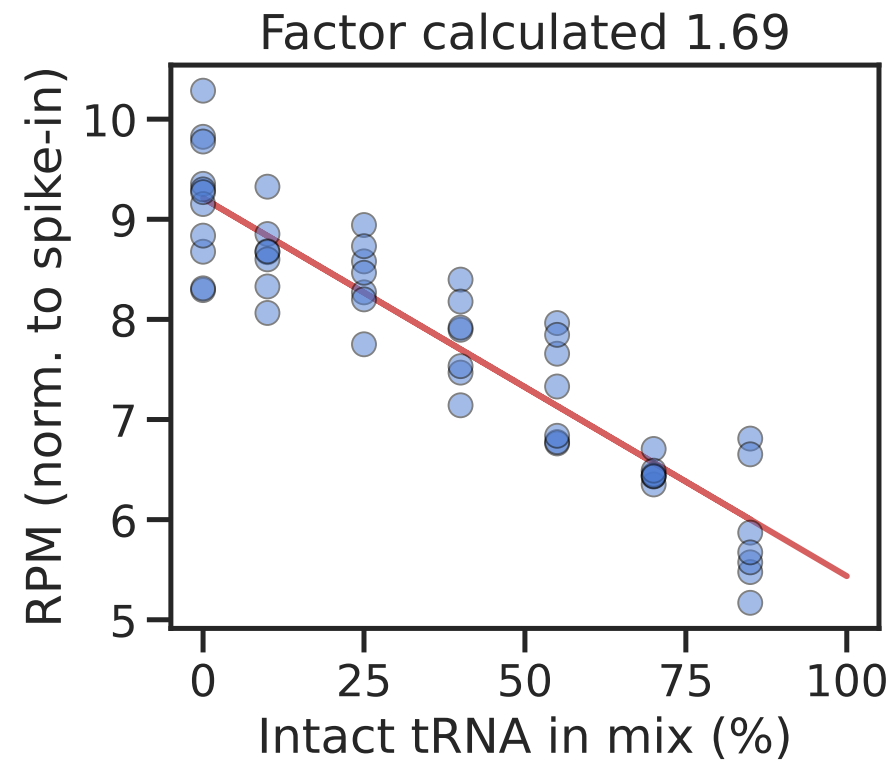
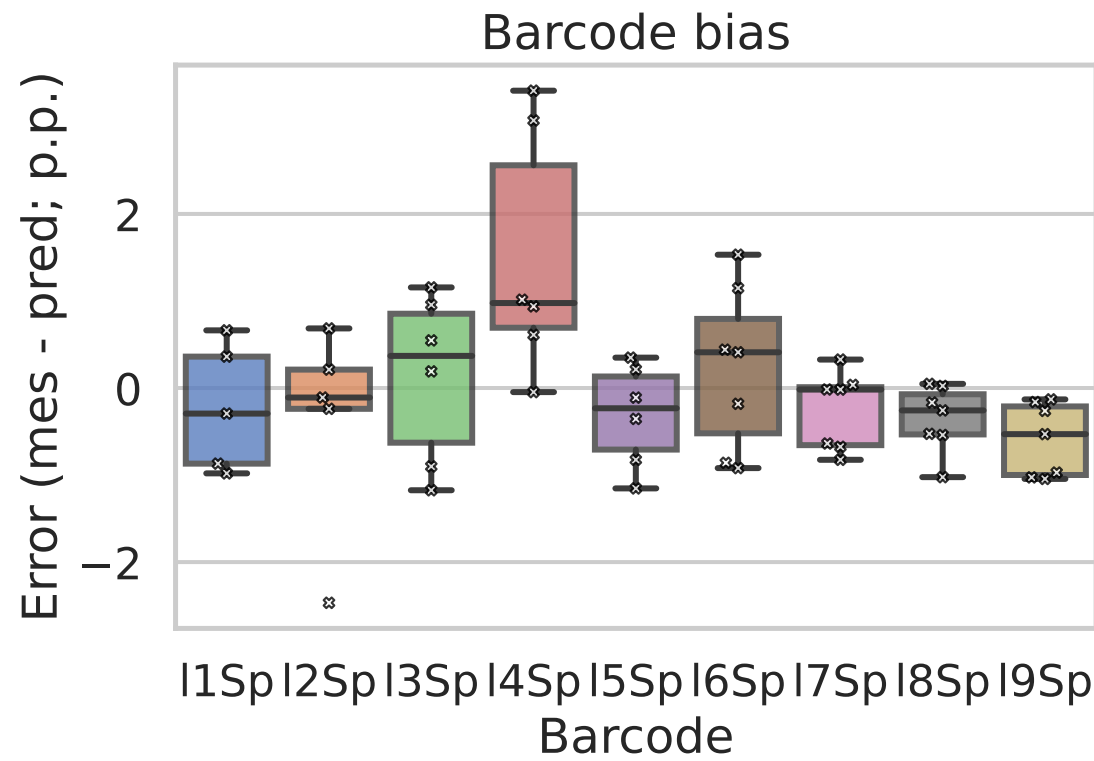
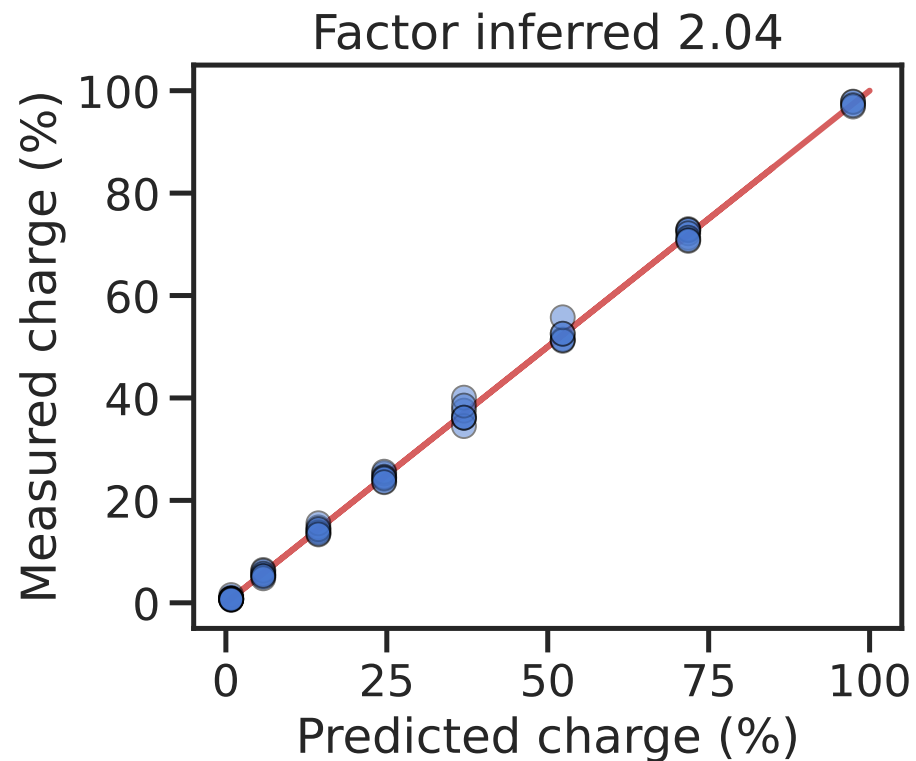
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Phe-GAA-2-1



Ile-TAT-1-1



Ala-AGC-1-1

