



**Supplementary Figure. Far UV Circular Dichroism Scans and Thermal Unfolding Curves for G $\beta$ 1-WT and Mutant Variants.** (a) The CD spectra of the nine variants are nearly identical to that of the wild-type protein. ■ MonB-WT, ■ MonB-A45V, ■ G $\beta$ 1-Y45A, ■ G $\beta$ 1-W43A, ■ G $\beta$ 1-W43Y, ■ MonB-ORDES, ■ G $\beta$ 1-W43V, ■ MonB-A45Y, ■ G $\beta$ 1-WT, ■ MonA (data not normalized). (b) Thermal denaturation of all variants monitored by CD at 218 nm (normalized as described in Supplementary Methods).

## **Supplementary Material**

### **Oligonucleotide Sequences**

#### **Construction of the Chimeric Construct Plasmid**

The gene for RNAP- $\alpha$  was amplified by PCR from the pTRG plasmid and cloned into the pBT vector downstream of  $\lambda$ cI via EcoRI/BamHI (oligonucleotides: 5'-GCGTCTG AATTCAT GCAGGGTTCTGTGACAGAGTTTCT-3' and 5'-GAATTATAGATCCGGCCGCCTCTGGTT TCTCTTCTTT-3'). The genes for the G $\beta$ 1 variants were sub-cloned into the chimeric vector with the engineered restriction sites NotI/EcoRI (oligonucleotides: 5'-AAGAGGCGGCCGCAT CTACTACTTACAAATTAATCCTTAA-3' and 5'-GGTGGTGATTCCCTTCAGTAACTGTA AAGGTCTTAGT-3').

#### **Creation of the G $\beta$ 1 Mutant Variants**

Point mutants of G $\beta$ 1-WT and MonB were produced using the QuikChange® method with the following oligonucleotides:

ProG W43Y F (5'GACAACGGTGTGACGGTGAATATACTTACGACGATGCGACTAAG)

ProG W43Y R (5'CTTAGTCGCATCGTCGTAAGTATATTCACCG TCAACACCGTTGTC)

ProG W43V F (5'GACAACGGTGTGACGGTGAAGTACTT ACGACGATGCGACTAAG)

ProG W43V R (5'CTTGTCGCATCGTCGTAAGTCACTT CACCGTCAACACCGTTGTC)

ProG W43A F (5'GACAACGGTGTGACGGTGAAGC GACTTACGACGATGCGACTAAG)

PROG W43A R (5'CTTAGTCGCATCGTCGTAA GTCGCTTCACCGTCAACACCGTTGTC)

MONB A45Y F (5'TTAAGGGTGAATGGA CAGTAGATGAAGCGACCAAGAC)

MONB A45Y R (5'GTCTTGGTCGCTTCATCTAC TGTCCATTCACCCTTAA)

MONB A45V F (5'TTAAGGGTGAATGGACATACGATGAAGCGACCAAGAC)

MONB A45V R (5'GTCTTGGTCGCTTCATCGTATGTCCATT CACCCTTAA).

### **Total Gene Synthesis via Recursive PCR**

Synthetic DNA oligonucleotides were used for recursive PCR synthesis of the genes for the MonB-ORDES variant and the genes in the randomized library.

1(B1+LIB): (5'GCGGCCGCATCTACCTATAAGCTGATTCTGAATGGCAAGACCCTGAA  
AGGTGAAACCACGACCGAA)

2(B2): (5' CCGCTGCATACTGTGCAAATACATCCTTTGCTGTTGCTTTGTCCACTGCTTC  
GGTCGTGGTTTCACCTTT)

3 (B3+LIB): (5'-TATTTGCACAGTATGCAGCGGATAACGGTGTTAAGGGTGAATGG)

4 (B4): (5'-GGAATTCCCTTCGGTCACGGTGAACGTCTTGGTCGCTTCGTCGAATGTCC  
ATT CACCCTTAACACCGTT)

5 (B\_PRIMER\_F) (5'GCGCACTATCGCGGCCGCATCTAC)

6 (B\_PRIMER\_R) (5'GCGCACGCTCGTGGAATTCCCTTCG)

2' (B\_LIB2): (5'CCGCTGCATACTGTGCAAATACATCXXXTGCTGTTGCXXXGTCCACT  
GCTTCGGTCGTGGTTTCACCTTT)

4' (B\_LIB4): (5'GGAATTCCCTTCGGTCACGGTGAACGTCTTGGTCGCTTCGTCXXXTG  
TCCATTCACCCTTAACACCGTT)

### **Protein Expression and Purification**

For protein expression the genes for the G $\beta$ 1 variants were sub-cloned into pET-21a (Novagen) using the restriction enzymes NdeI and EcoRI (oligonucleotides: 5'-GTC CGC GGT CAT ATG ACT ACT TAC AAA-3' and 5'-GGC GCA GAT GAA TTC TTA TTC AGT AAC TGT AAA-3') and transformed into BL21(DE3).