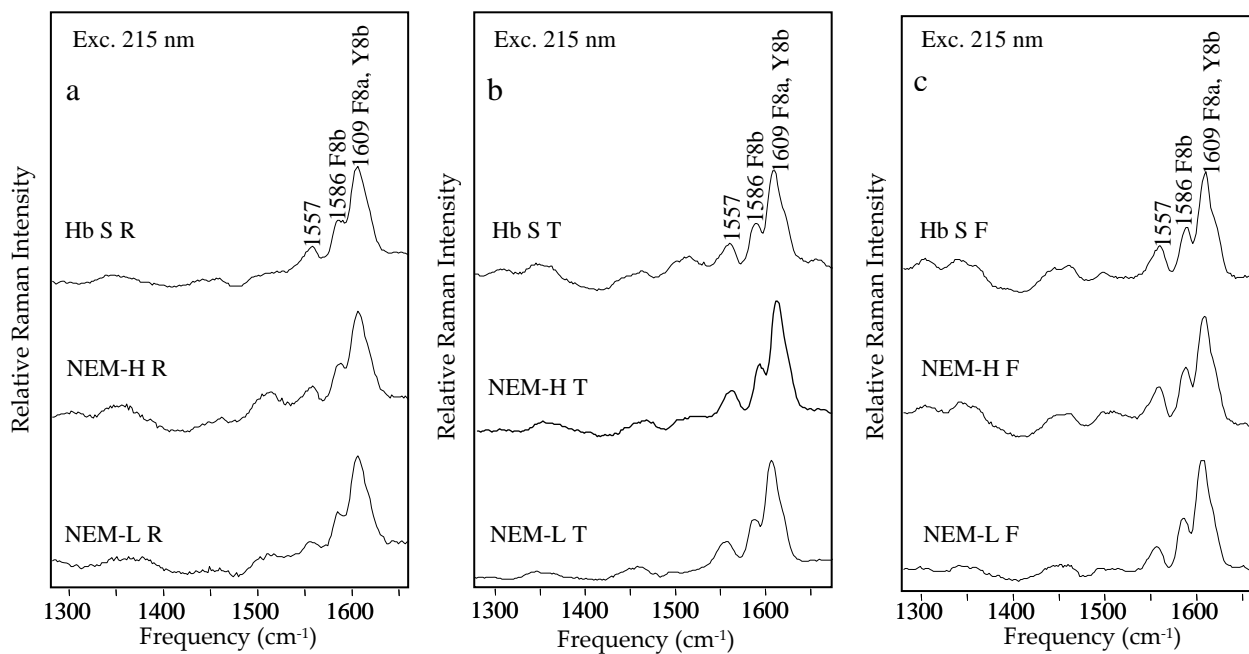
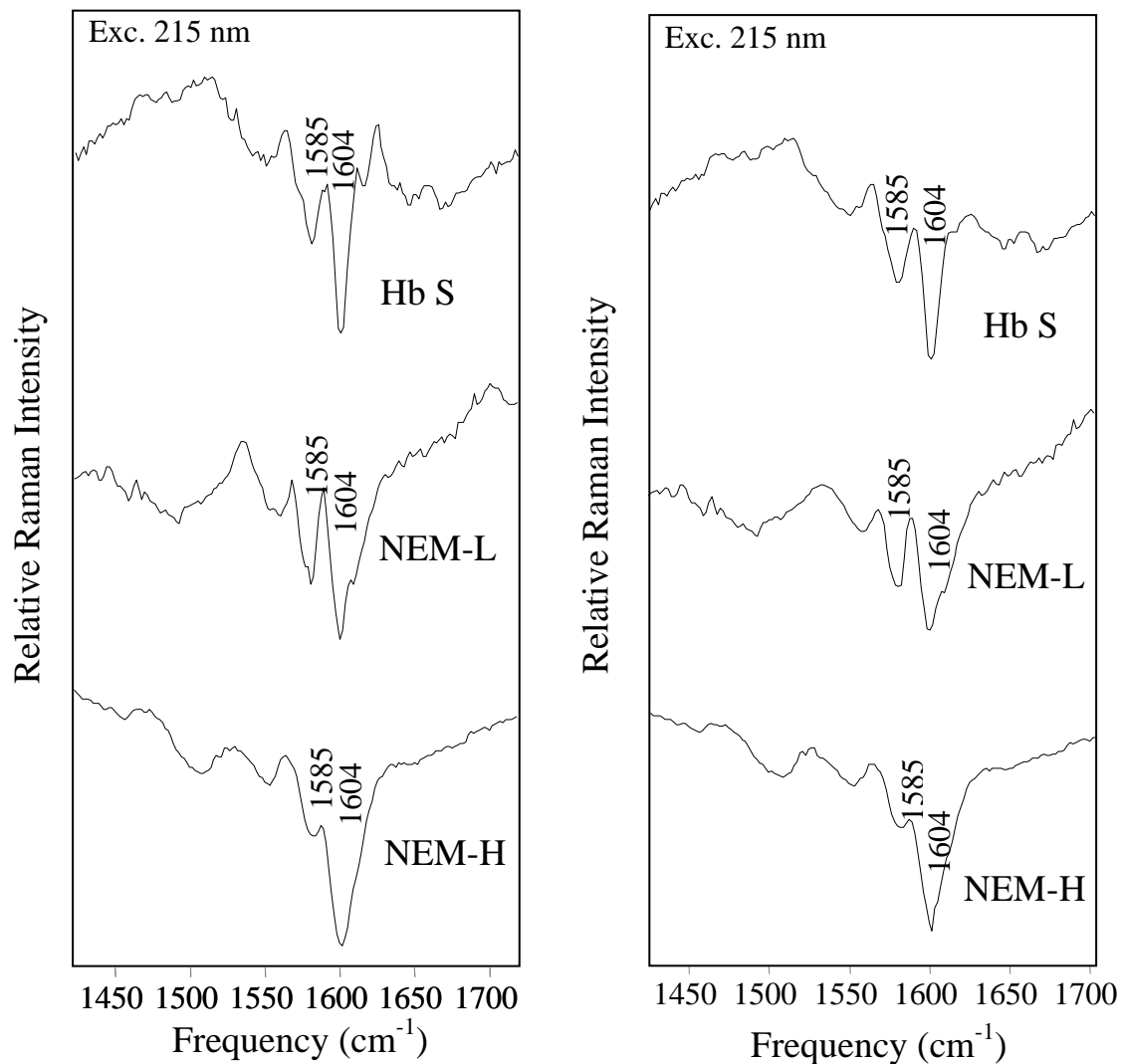


Supplementary Figures



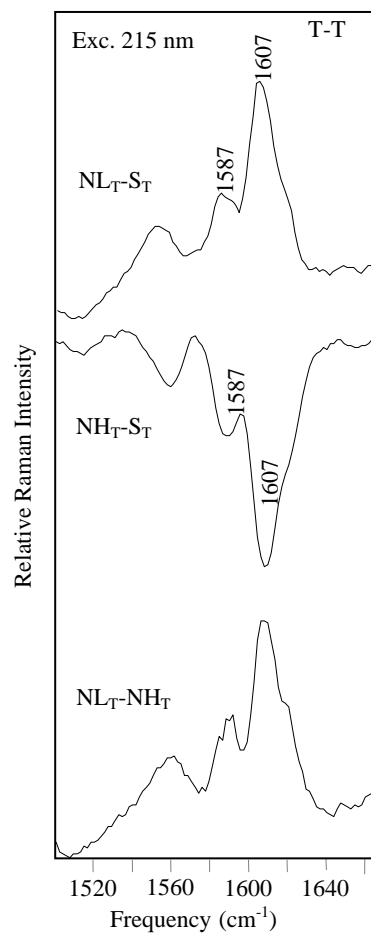
Supplementary Figure 1

Supplementary Figure 1: 215 nm-excited UVRR spectra. R- (a), T- (b), and Fiber-state (c), spectra of Hb S (top), NEM-H Hb (center) and NEM-L Hb (bottom) obtained with 215 nm excitation. Samples were prepared as described in figures 5 and 6. Depicted spectra result from 3 hours of data acquisition.



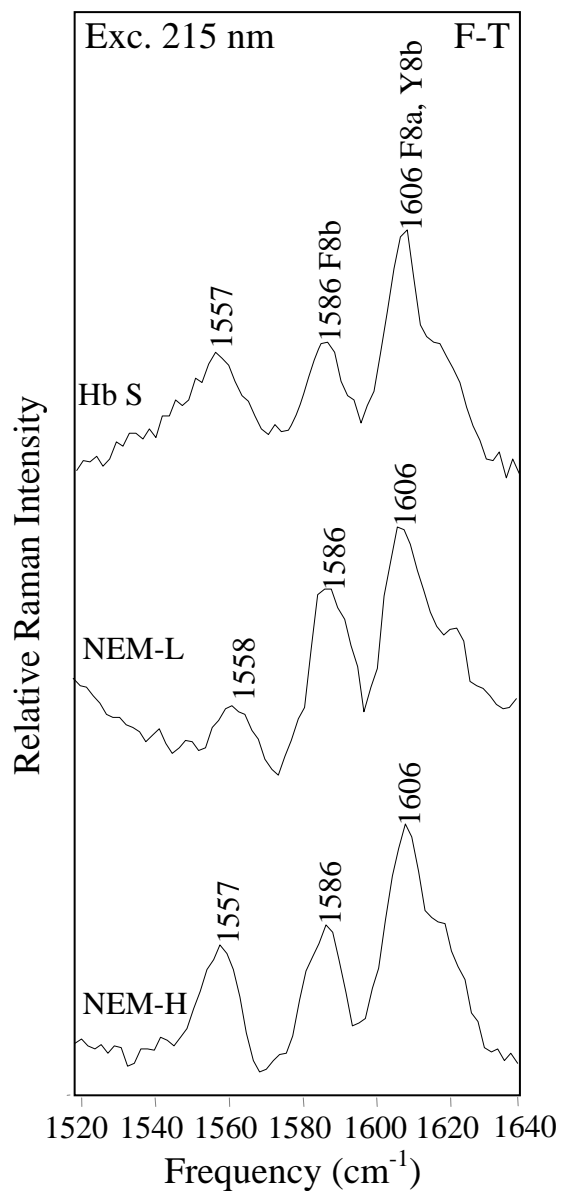
Supplementary Figure 2.

Supplementary Figure 2: T-R difference spectra of Hb S (top), NEM-H Hb (center) and NEM-L Hb (bottom), obtained with 215 nm excitation, in high (left) and low (right) potassium phosphate buffer.. Low phosphate samples were 0.25 mM Hb, in a 0.1 potassium phosphate buffer, pH 7.1, and high phosphate samples were 1.0 mM Hb in a 1.0 mM potassium phosphate buffer, pH 7.1. Samples were prepared as described in Materials and Methods.



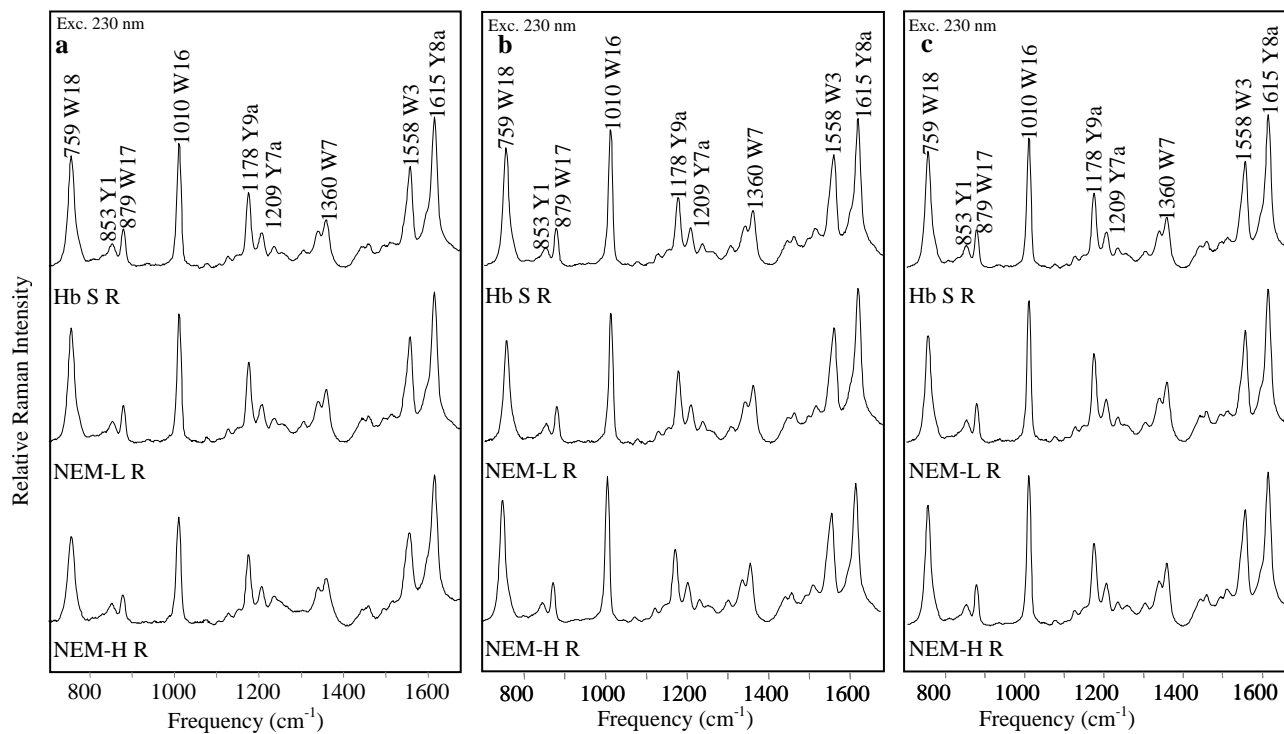
Supplementary Figure 3

Supplementary Figure 3: T-T difference spectra of Hb S (top), NEM-H Hb (center) and NEM-L Hb (bottom), obtained with 215 nm excitation. Samples were 0.25 mM Hb, in a 0.1 potassium phosphate buffer, pH 7.1. Samples were prepared as described in Materials and Methods.



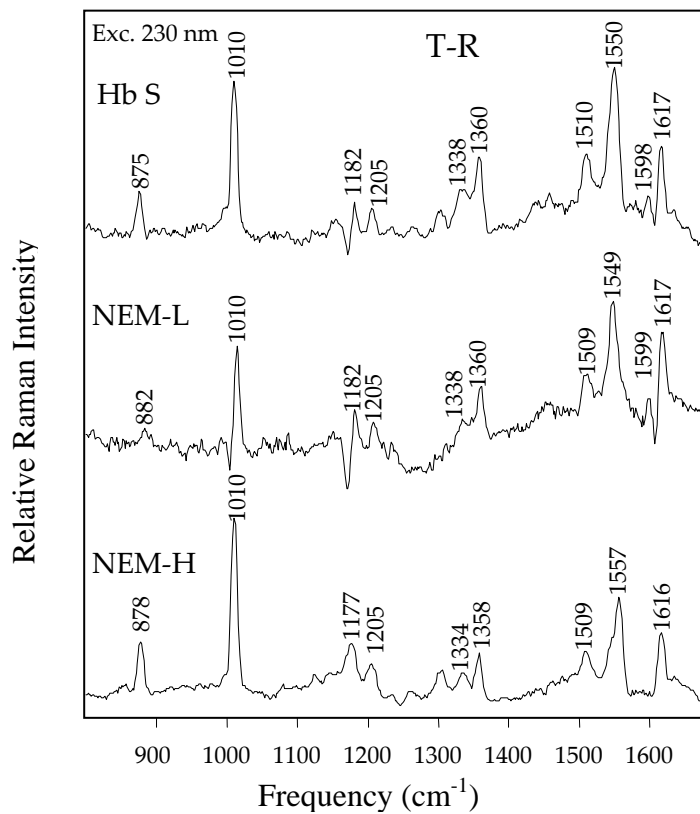
Supplementary Figure 4

Supplementary Figure 4: F-T difference spectra of Hb S (top), NEM-H Hb (center) and NEM-L Hb (bottom), obtained with 215 nm excitation. T-state samples were 0.25 mM Hb, in a 0.1 potassium phosphate buffer, pH 7.1, and fiber samples were 3.0 mM Hb in a 0.1 M potassium phosphate buffer, pH 7.1. Samples were prepared as described in Materials and Methods.



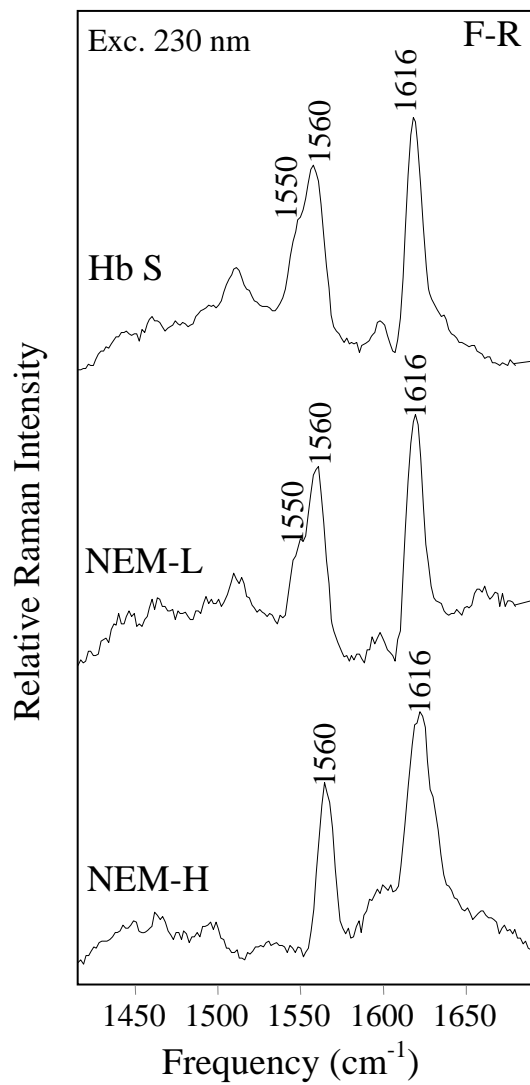
Supplementary Figure 5

Supplementary Figure 5: 230 nm excited UVRR spectra. R- (a), T- (b), and Fiber-state (c) spectra of Hb S (top), NEM-H Hb (center) and NEM-L Hb (bottom) obtained with 230 nm excitation. Samples were prepared as described in figures 5 and 6. Depicted spectra result from 3 hours of data acquisition.



Supplementary Figure 6

Supplementary Figure 6: T-R difference spectra of Hb S (top), NEM-L (center) and NEM-H (bottom) obtained with 230 nm excitation. Samples were 0.25 mM Hb in a 0.1 M potassium phosphate buffer, pH 7.1.



Supplementary Figure 7

Supplementary Figure 7: F-R difference spectra of Hb S (top), NEM-L (center) and NEM-H (bottom) obtained with 230 nm excitation. R-state samples were 0.25 mM Hb in a 0.1 M potassium phosphate buffer, pH 7.1, and fiber samples were 3.0 mM Hb in a 0.1 M potassium phosphate buffer, pH 7.1.

Supplementary Table 1

Supplementary Table 1: List of sequence of all Cys containing labeled peptides associated with the β 93 and α 104 Cys residues, observed by reverse phase LC mass spectrometry.

α 104

VNFKLLSHCLLV
VNFKLLSHCLLVT
VNFKLLSHCLLVTLA
VNFKLLSHCLLVTLAA
VNFKLLSHCLLVTLAAHLPAAE
AHKLRVDPVNFKLLSHCLL
FKLLSHCLL
FKLLSHCLLV
FKLLSHCLLVTLA
FKLLSHCLLVTLAA
FKLLSHCLLVTLAAH
FKLLSHCLLVTLAAHLPAAEFT
FKLLSHCLLVTLAAHLPAAEFTPAV
KLLSHCLL
KLLSHCLLVTLA
KLLSHCLLVTLAA
KLLSHCLLVTLAAH
KLLSHCLLVTLAAHLPAAEFT
KLLSHCLLVTL
KLRVDPVNFKLLSHCLL
KLRVDPVNFKLLSHCLLV
KLRVDPVNFKLLSHCLLVTL
KLRVDPVNFKLLSHCLLVTLA
LHAHKLRVDPVNFKLLSHCLL
LLSHCLL
LLSHCLLV
LLSHCLLVTL
LLSHCLLVTLA
LLSHCLLVTLAAH
LLSHCLLVTLAAHLPAAE
LLSHCLLVTLAAHLPAAEFT
LLSHCLLVTLAAHLPAAEFTPAV
LRVDPVNFKLLSHCLL
LRVDPVNFKLLSHCLLV
LRVDPVNFKLLSHCLLVTL
LSHCLLVTLAAH
RVDPVNFKLLSHCLLV
RVDPVNFKLLSHCLLVTL

RVDPVNFKLLSHCLLVTLA
RVDPVNFKLLSHCLLVTLAAHLP
AE
RVDPVNFKLLSHCLL
RVDPVNFKLLSHCLLV
RVDPVNFKLLSHCLLVT
RVDPVNFKLLSHCLLVTLAA
RVDPVNFKLLSHCLLVTLAAH
VDPVNFKLLSHCLL
VDPVNFKLLSHCLLV
VDPVNFKLLSHCLLVT
VDPVNFKLLSHCLLVTLA
VDPVNFKLLSHCLLVTLAA
VNFKLLSHCLLV
VNFKLLSHCLLVT
VNFKLLSHCLLVTLA
VNFKLLSHCLLVTLAA
VNFKLLSHCLLVTLAAHLP
AE

β93

AFSDGLAHLDNLKGTFATLSELHCDKL
AHLDNLKGTFATLSELHCDKLH
AHLDNLKGTFATLSELHCDKLHVDPENFRL
AHLDNLKGTFATLSELHCDKLHVDPENFRLL
ATLSELHCDKLH
ATLSELHCDKLHVD
ATLSELHCDKLHVDPENFRLL
ATLSELHCDKLHVDPENFRLLGNVL
ATLSELHCDKL
ATLSELHCDKLHVDPENFR
ATLSELHCDKLHVDPENFRLLGNVLV
ATLSELHCDKLHVDPENFRLLGNVLVCVL
FATLSELHCDKL
FATLSELHCDKLH
FATLSELHCDKLHVD
FATLSELHCDKLHVDPENFR
FATLSELHCDKLHVDPENFRL
FATLSELHCDKLHVDPENFRLLG
KGT
FATLSELHCDKL
KGT
FATLSELHCDKLH
KGT
FATLSELHCDKLHVD
KGT
FATLSELHCDKLHVDPENFRL
KGT
FATLSELHCDKLHVDPENFRLLGNVLV
LDNLKGTFATLSELHCDKLHVDPENFR
LDNLKGTFATLSELHCDKL
LGA
FSDGLAHLDNLKGTFATLSELHCDKL
LHCDKLH

LHCDKLHVD
LHCDKLHVDPENFRL
LHCDKLHVDPENFRLL
LHCDKLHVDPENFRLLG
LHCDKLHVDPENFRLLGNVLV
LHCDKLHVDPENFRLLGNVLVCVL
LHCDKLHVDPENFR
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