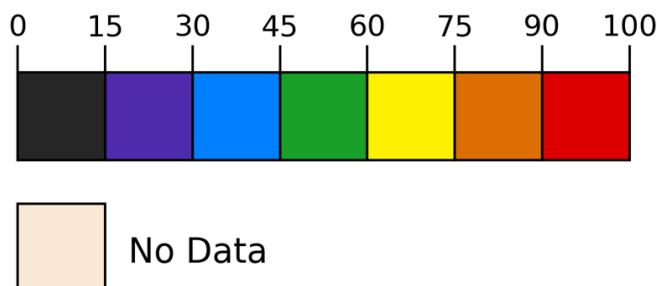


Creating Custom Color Schemes

By inputting the lower cutoff for a color to be applied to peptides, as well as the color, figure colors can be completely customizable. The text color of any text that appears over a peptide can also be modulated. Here are some examples of the inputs that generated the default coloring schemes, and the associated legend created dynamically by the software attached.

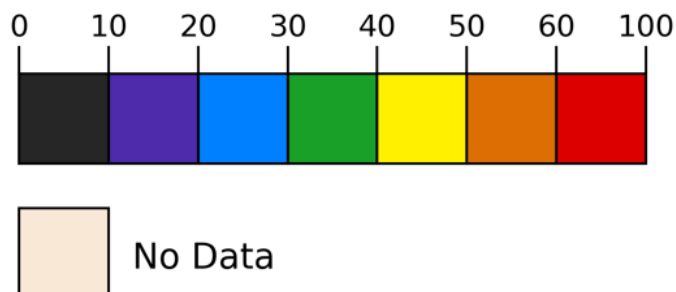
1) Default Corrected Uptake Colors - “uptake_cor_default.json”

| Create Custom Colors for All Uptake Maps | | |
|---|-------------------|-------------------------------------|
| Enter RFU as a decimal, with the highest exchanging color first | | |
| | Hexadecimal Color | White Text |
| If RFU > 0.9 : | DC0000 | <input checked="" type="checkbox"/> |
| If RFU > 0.75 : | DC6E03 | <input checked="" type="checkbox"/> |
| If RFU > 0.6 : | FFF000 | <input type="checkbox"/> |
| If RFU > 0.45 : | 19A028 | <input checked="" type="checkbox"/> |
| If RFU > 0.3 : | 007FFF | <input checked="" type="checkbox"/> |
| If RFU > 0.15 : | 4D2BAB | <input checked="" type="checkbox"/> |
| If RFU > : | | <input checked="" type="checkbox"/> |
| If RFU > : | | <input checked="" type="checkbox"/> |
| If RFU > : | | <input checked="" type="checkbox"/> |
| If RFU > 0 : | 262626 | <input checked="" type="checkbox"/> |
| If RFU < 0 : | 262626 | <input checked="" type="checkbox"/> |
| If RFU = 0 : | F2F2F2 | <input checked="" type="checkbox"/> |
| If Peptide Absent : | FAE8D7 | <input checked="" type="checkbox"/> |



Note: Not all rows need to be filled. All text displayed on the yellow color here (FFF000) will appear as black for maximal visibility.

2) Default Uncorrected Uptake Colors – “uptake_uncor_default.json”



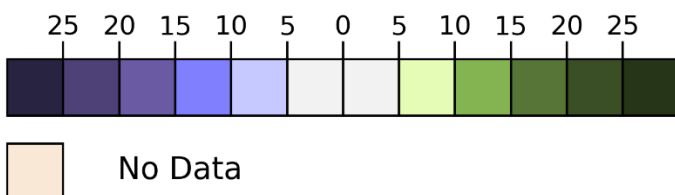
HDXWizard

Create Custom Colors for All Uptake Maps

Enter RFU as a decimal, with the highest exchanging color first

| | Hexadecimal Color | White Text |
|---------------------|-------------------|-------------------------------------|
| If RFU > 0.6 : | DC0000 | <input checked="" type="checkbox"/> |
| If RFU > 0.5 : | DC6E03 | <input checked="" type="checkbox"/> |
| If RFU > 0.4 : | FFF000 | <input type="checkbox"/> |
| If RFU > 0.3 : | 19A028 | <input checked="" type="checkbox"/> |
| If RFU > 0.2 : | 007FFF | <input checked="" type="checkbox"/> |
| If RFU > 0.1 : | 4D2BAB | <input checked="" type="checkbox"/> |
| If RFU > : | | <input checked="" type="checkbox"/> |
| If RFU > : | | <input checked="" type="checkbox"/> |
| If RFU > : | | <input checked="" type="checkbox"/> |
| If RFU > 0 : | 262626 | <input checked="" type="checkbox"/> |
| If RFU < 0 : | 262626 | <input checked="" type="checkbox"/> |
| If RFU = 0 : | F2F2F2 | <input checked="" type="checkbox"/> |
| If Peptide Absent : | FAE8D7 | <input checked="" type="checkbox"/> |

3) Default Experimental (maxD) Difference Colors – “Exp_dif_default.json”



Create Custom Colors for All Difference Maps

Is this difference in Daltons (Theoretical) or RFU (Experimental)

Enter Difference with the highest absolute value differences first. For RFU enter as a decimal

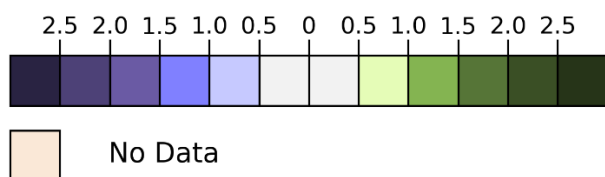
| Protection | | | Deprotection | | |
|---------------------|-------------------------------------|-------------------------------------|-----------------|-------------------------------------|-------------------------------------|
| | Hexadecimal Color | White Text | | Hexadecimal Color | White Text |
| If dif > 0.25 : | <input type="text" value="292342"/> | <input checked="" type="checkbox"/> | If dif > 0.25 : | <input type="text" value="263418"/> | <input checked="" type="checkbox"/> |
| If dif > 0.2 : | <input type="text" value="4D4177"/> | <input checked="" type="checkbox"/> | If dif > 0.2 : | <input type="text" value="3A4F25"/> | <input checked="" type="checkbox"/> |
| If dif > 0.15 : | <input type="text" value="6A5AA4"/> | <input checked="" type="checkbox"/> | If dif > 0.15 : | <input type="text" value="567537"/> | <input checked="" type="checkbox"/> |
| If dif > 0.1 : | <input type="text" value="8080FF"/> | <input checked="" type="checkbox"/> | If dif > 0.1 : | <input type="text" value="84B451"/> | <input checked="" type="checkbox"/> |
| If dif > 0.05 : | <input type="text" value="C6C9FF"/> | <input type="checkbox"/> | If dif > 0.05 : | <input type="text" value="E5FCB7"/> | <input type="checkbox"/> |
| If dif > 0 : | <input type="text" value="F2F2F2"/> | <input type="checkbox"/> | If dif > 0 : | <input type="text" value="F2F2F2"/> | <input type="checkbox"/> |
| If dif = 0 : | <input type="text" value="F2F2F2"/> | <input type="checkbox"/> | | | |
| If Peptide Absent : | <input type="text" value="FAE8D7"/> | <input checked="" type="checkbox"/> | | | |

[See examples](#)

[Save Colors](#)

Note: Differences here are calculated by percent difference between maxD corrected RFUs.

4) Default Theoretical (absolute difference) Difference Colors - "Theo_dif_default.json"



Create Custom Colors for All Difference Maps

Is this difference in Daltons (Theoretical) or RFU (Experimental)

Enter Difference with the highest absolute value differences first. For RFU enter as a decimal

| Protection | | | Deprotection | | |
|---------------------|-------------------|-------------------------------------|----------------|-------------------|-------------------------------------|
| | Hexadecimal Color | White Text | | Hexadecimal Color | White Text |
| If dif > 2.5 : | 292342 | <input checked="" type="checkbox"/> | If dif > 2.5 : | 263418 | <input checked="" type="checkbox"/> |
| If dif > 2 : | 4D4177 | <input checked="" type="checkbox"/> | If dif > 2 : | 3A4F25 | <input checked="" type="checkbox"/> |
| If dif > 1.5 : | 6A5AA4 | <input checked="" type="checkbox"/> | If dif > 1.5 : | 567537 | <input checked="" type="checkbox"/> |
| If dif > 1 : | 8080FF | <input checked="" type="checkbox"/> | If dif > 1 : | 84B451 | <input checked="" type="checkbox"/> |
| If dif > 0.5 : | C6C9FF | <input type="checkbox"/> | If dif > 0.5 : | E5FCB7 | <input type="checkbox"/> |
| If dif > 0 : | F2F2F2 | <input type="checkbox"/> | If dif > 0 : | F2F2F2 | <input type="checkbox"/> |
| If dif = 0 : | F2F2F2 | <input type="checkbox"/> | | | |
| If Peptide Absent : | FAE8D7 | <input checked="" type="checkbox"/> | | | |