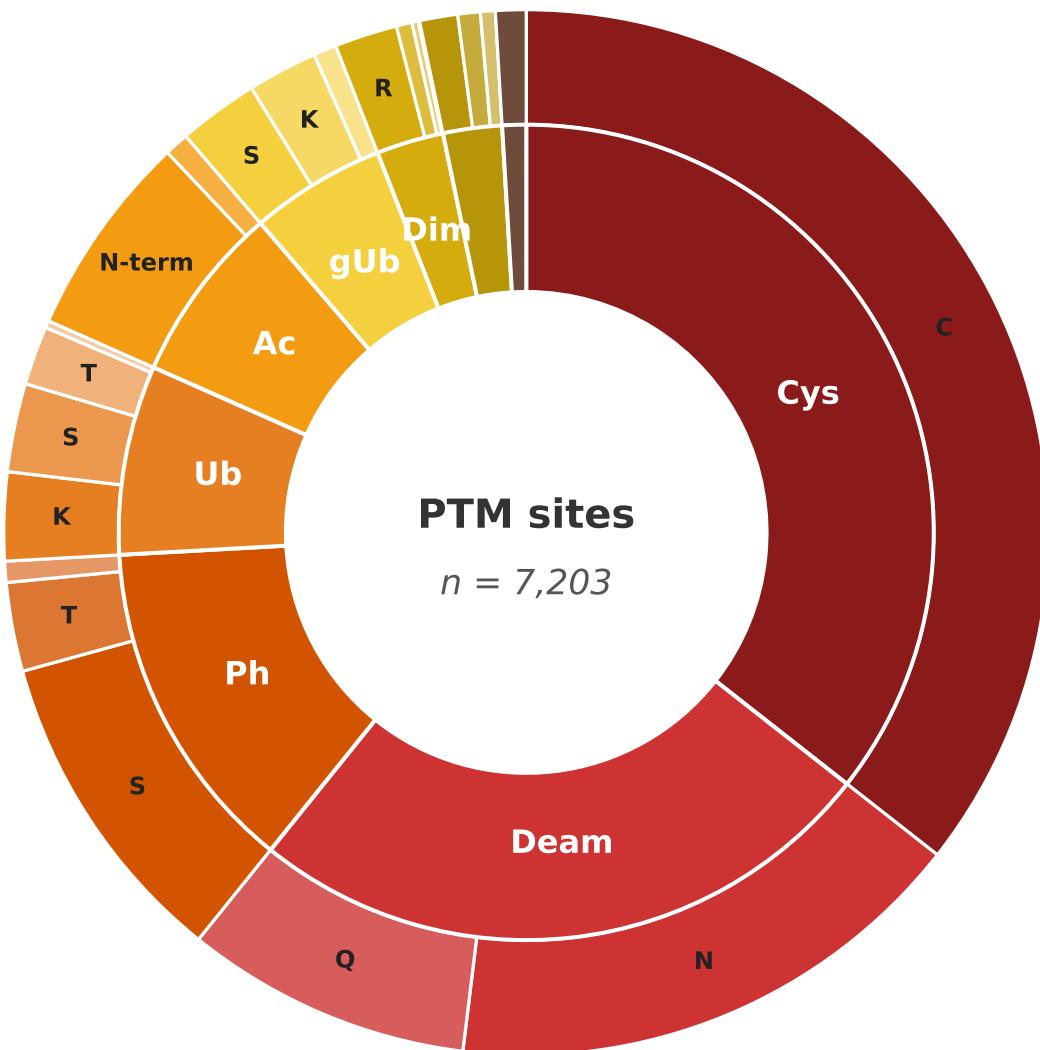


# HLA-I PTM Distribution by Modification Type and Residue



PTM Types	
Cysteinylation	(2,563, 35.6%)
Deamidation	(1,816, 25.2%)
Phosphorylation	(960, 13.3%)
Ubiquitination (GG)	(541, 7.5%)
Acetylation	(509, 7.1%)
Ubiquitination (G)	(386, 5.4%)
Dimethylation	(192, 2.7%)
Methylation	(168, 2.3%)
SUMOylation	(68, 0.9%)