ZipChip Charge Variant Analysis

Many current mAb characterization techniques either provide high-resolution charge variant separations or mass spec data. Not both. This makes identification of mAb variants difficult.

No more either/or

ZipChip[®] can perform high-resolution charge variant separations of intact antibodies backed by the power of Native-MS characterization. In a single run you can get mass information, assess charge heterogeneity, characterize glycoforms, identify basic and acidic variants, and more. The added layer of mass spec information means you can finally pinpoint the identity of each peak.

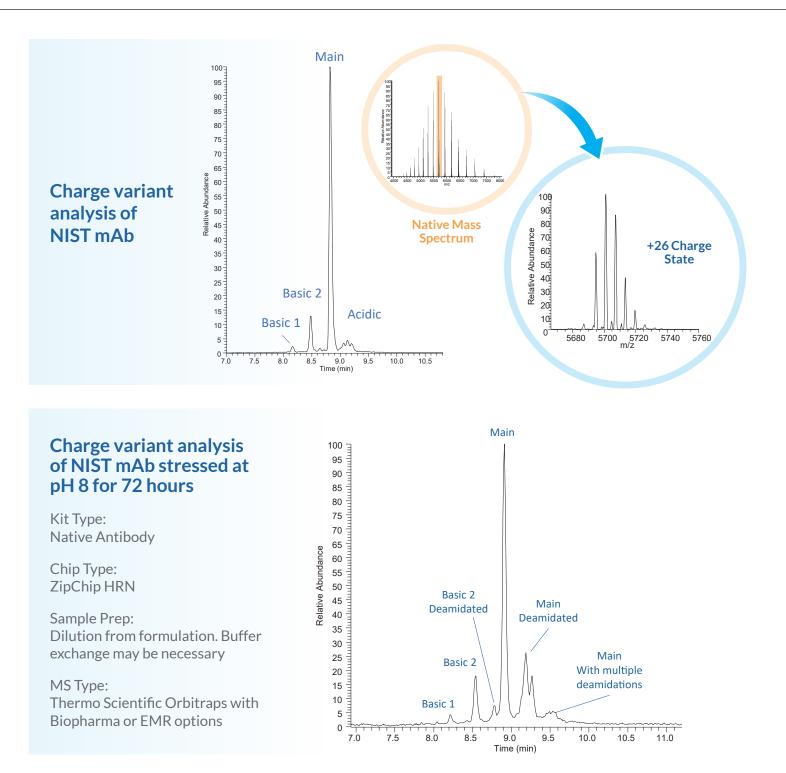
Get the unfair advantage

When compared to traditional methods for charge variant analysis of antibodies, ZipChip separates more and allows mass spec identification for each charge variant.

| | Charge Variant Separation | MS Compatible | Charge Variant Masses | Charge Variant ID |
|-------------------------|------------------------------|---------------|--------------------------|-------------------|
| ZipChip | **** | **** | **** | **** |
| lsoelectric focusing | **** | × | × | ** |
| CE (UV) | **** | × | × | ** |
| RPLC-MS | * | **** | ** | ** |



ZipChip





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