COMPLETE REPORTING OF ALL PEAKS (CRAP)

is the most commonly used protocol in metabolomics.

ARE YOU STILL PRODUCING CRAP DATA?

(Yet you know they are mostly artifacts, adducts, and fragments, and all the peaks are suppressed?)

The IROA TruQuant protocol and ClusterFinder software ensure data quality

- Provides quantitation and identification (with built-in libraries) for hundreds of IROA and non-IROA peaks in your sample
- Automatically corrects for most variances, including instrumental, matrix effect and sample preparation
- Sample-to-sample normalization
- Corrects for all ion losses
- Assures quality, reproducible data
- Economical less than the cost of the plasticware used in preparing a sample

Why not make your data worth working up?
PUT THE CRAP BEHIND YOU AND MOVE TO IROA-DRIVEN ACCURATE DATA

IROA TECHNOLOGIES

...making metabolic profiling easy®