

Laboratory for Proteomics
ERA Chair project
" VetMedZg"

PRICE IN EURO:	FVM	Academia	Industry
SAMPLE PREPARATION			
1. Concentration determination	15	23	30
2. Depletion			
2.1. Albumin depletion	15	23	30
2.2. IgG depletion	15	23	30
2.3. ProteoMiner	80	120	160
3. Digestion			
3.1. In solution	15	23	30
3.2. In gel	15	23	30
3.3. FASP	30	45	60
4. Desalting/ detergent removal	25	38	50
5. Desalting/ protein concentration	15	23	30
PROTEIN ID			
1. From gel spots/bands			
1.1. In gel digestion	15	23	30
1.2. Min 60 min runtime	60	90	120
2. Shotgun			
2.1. FASP	30	45	60
2.2. Min 150 min runtime	90	133	180
PROTEIN RELATIVE QUANTIFICATION (TMT)*			
1. In solution digestion/precipitation	20	30	40
2. TMT labeling	310/60**	355/80	400/100
3. Min 150 min runtime	90	135	180
DATA ANALYSIS			
1. Proteome Discoverer	15	23	30

***For two samples (minimum number)**

**** The user is buying chemicals for the relative quantification of the protein by the TMT method**

Basic price per sample/EUR

For samples in MS non-compatible buffers, FASP or desalting is needed!
For higher number of samples, discount can be applied!

Example of analysis workflow

