

## N-Terminal Edman Sequencing

Priority  normal  high Internal sample number .....

### Client

Name\* ..... email\* .....

Institution\* .....

Address\* .....

Phone, fax\* .....

Cost model\*:  Inst. Biochemistry

Industry

other: .....

No of Cycles (min. 6 cycles)\* : .....

### Sample <sup>a</sup>:

Protein  Peptide

Description : .....

Sequence ID : .....

Electronic sequence file  no  yes : .....

Amount\* : ..... Purity \* : .....

Sample preparation : .....

known contaminants :  salt :

detergents : .....

known modifications

Radioactivity\*  no  yes : .....

dry soluble in  water  0.1% TFA  other : .....

dissolved in  water  0.1% TFA  other : .....

blotted (PVDF)

Storage\*  -80°C  -20°C  4-8°C  RT

Signature client .....

Date: .....

Fax signed form to: 0641-99-47489

\* required

<sup>a</sup> Detailed information about the sample is a prerequisite for successful analysis. Clients are strongly encouraged to discuss sample preparation with the core facility team: [proteomics@biochemie.med.uni-giessen.de](mailto:proteomics@biochemie.med.uni-giessen.de)