

PROTEIN/PEPTIDE SEQUENCE ANALYSIS SUBMISSION FORM

RESE	ARCHER INFORMATION	BILLING INFORMATION	
Date: Principal Investigator: Department/Company:		P.O./MC/VISA #: Exp. Date (if applicable):	
		Phone:	Fax:
SHIPPING ADDRESS:		BILLING ADDRESS:	
		-	
	Sample Inf		
	Please provide as much	information as possible:	
1.	SAMPLE AOUNT: moles; or micrograms		
	A. Recommend minimum amount of samples is 100 pMoles. Example: 100 pmoles of a protein (M.W. 10,000) would weigh 1.0 µg		
			B. If less than 100 pmoles of protein sample, there are NO GUARANTEES of adequate sequencing results.
	2.	IS SAMPLE in SOLUTION or DRY	
2.	A. If sample is in solution, what is the sample stored is	n?	
	11. If sample is in solution, what is the sample stored in:		
3.	METHOD(s) USED TO OBTAIN THE SAMPLE:		
	A. For samples electroblotted to PVDF: What membrane was used?		
	Immobilon-P (.45 micron) (Millipore)	Problot (.1 micron) (ABI)	
	Westran (.45 micron) (Schleicher & Schuell	Trans-Blot (.1 micron) (Biorad)	
	Immobilon-PSQ (.1 micron) (Millipore)	Fluoratrans (.1 micron) Pall Corp)	
	B. It is possible to sequence directly from a membrane	:: PVDF & Immobilon	
4.	ESTIMATED PURITY OF THE SAMPLE:		
	A. Method(s) used to obtain Purity:		
	B. If your sample was collected on a HPLC, please at and wavelength	ttach the chromatogram with gradient, solvents, column	