

Internal Application Form

Host institution	
Kind of occupation	
Earliest entry date	

ESR choice (ESR ID No.)	1st choice ESR ID No. _____	2nd choice ESR ID No. _____	3rd choice ESR ID No. _____
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PERSONAL DATA					
Surname				First name(s)	
Title		Date of Birth	<small>(dd.mm.yyyy)</small>	Sex	<input type="checkbox"/> Female <input type="checkbox"/> Male
Citizenship			Zip code / Place		
Street/Number/Top					
Social security no. <small>(where applicable)</small>				Phone	
Marital state	<input type="checkbox"/> single <input type="checkbox"/> married <input type="checkbox"/> partnership				

Is there any disability / reduction in earning capacity?	
<input type="checkbox"/> yes, degree of disability: _____	<input type="checkbox"/> no

EDUCATION			
School / university	Subject	From – to	Completed
			<input type="checkbox"/> yes <input type="checkbox"/> no
			<input type="checkbox"/> yes <input type="checkbox"/> no
			<input type="checkbox"/> yes <input type="checkbox"/> no
			<input type="checkbox"/> yes <input type="checkbox"/> no
			<input type="checkbox"/> yes <input type="checkbox"/> no

PREVIOUS EMPLOYMENT (in the past 5 years)			
Employer	Address	From – to	Kind of occupation

Languages - please refer to EU Common Reference Levels: https://www.examenenglish.com/CEFR/cefr.php	English	A1	A2	B1	B2	C1	C2
		A1	A2	B1	B2	C1	C2
		A1	A2	B1	B2	C1	C2
		A1	A2	B1	B2	C1	C2

None

<p>Project-specific Skills</p> <p>NB: In the 2nd step of recruitment, proof of skills will be required!</p>	Knowledge of network theory	excellent	very good	good	basic	
	Analysis of data sets from proteomics & transcriptomics	excellent	very good	good	basic	
	Time series gene expression analysis experience	excellent	very good	good	basic	
	Knowledge of machine learning methods	excellent	very good	good	basic	
	Data modelling (ideally semantic web RDF)	excellent	very good	good	basic	
	Knowledge of mathematical modelling	excellent	very good	good	basic	
	Experimental experience within affinity proteomics or similar	excellent	very good	good	basic	
	Cell cultures	excellent	very good	good	basic	
	Generation of extracellular vesicles	excellent	very good	good	basic	
	Experience in cell biology / histology	excellent	very good	good	basic	
	Experience in image analysis	excellent	very good	good	basic	
	Experience in data analysis	excellent	very good	good	basic	
	Practical data transformation skills (e.g. ETL, changing data from one format to another)	excellent	very good	good	basic	
	Medical statistics	excellent	very good	good	basic	
	Statistics skills / statistics background	excellent	very good	good	basic	
	Scripting/Programming language development	excellent	very good	good	basic	
	Familiarity with statistical software Stata	excellent	very good	good	basic	
	Scripting in C/C++	excellent	very good	good	basic	
	Scripting in R	excellent	very good	good	basic	
	Scripting in Matlab	excellent	very good	good	basic	
	Scripting in Python	excellent	very good	good	basic	
	Computing skills	excellent	very good	good	basic	
	Analytical and problem-solving skills	excellent	very good	good	basic	
	Knowledge of privacy implications when using health data	excellent	very good	good	basic	
	Awareness of privacy preserving techniques (anonymization/pseudonymisation)	excellent	very good	good	basic	
	Use of health and omics related data for research	excellent	very good	good	basic	
	Knowledge of GDPR, international standards (e.g. ISO 27001), data subject consent	excellent	very good	good	basic	
Other:	excellent	very good	good	basic		
Other:	excellent	very good	good	basic		
Other:	excellent	very good	good	basic		

Comments	
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I hereby confirm the validity and correctness of the above information with my personal signature. I am aware that false statements may result in exclusion from the recruitment process.

Date		Signature	
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