



# GLOBAL MIGRATION FLOWS OF SCIENTISTS AND THE STATUS OF INTERNATIONAL SCIENTISTS IN GERMANY

Emilio Zagheni (Max Planck Institute for Demographic Research)

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Ebru Sanliturk



Maciej Danko



**Tom Theile** 



**MPIDR Summer Incubator 2023** 

### **EMMANUELLE CHARPENTIER - NOBEL PRIZE WINNER 2020**







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Cell Reports Report

#### Structural Basis for Recognizing Phosphoarginine and Evolving Residue-Specific Protein Phosphatases in Gram-Positive Bacteria

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http://dx.doi.org/10.1016/j.celreg 2013 p.5.023

#### SUMMARY

Many cellular pathways are regulated by the competing activity of protein kinases and phosphatases. The recent identification of arginine phosphorylation as a protein modification in bacteria prompted us to analyze the molecular basis of targeting phosphoarginine. In this work, we characterize an annotated

of specific client proteins. Based on their sequence, structure, and function, protein phosphatases are grouped into three main classes. Phosphatases acting on phospho-serine/three-nine (pSer, pTm) comprise the PPP (phospho protein phosphatase) and PPM (Mg<sup>2</sup>/Mm<sup>2</sup>-dependent protein phosphatase) and PPM (Mg<sup>2</sup>/Mm<sup>2</sup>-dependent protein phosphatase) families, whereas enzymes acting on phospho-tyrosine (pTyr) constitute the protein tyrosine phosphatase (PTP) superfamily (Barford et al., 1998; Stoker, 2005). In addition, specialized protein phosphatases and se phospho-appadite, phosphatase







GENOME EDITING

## The new frontier of genome engineering with CRISPR-Cas9

Jennifer A. Doudna\* and Emmanuelle Charpentier\*

Structural Basis for Recognizing Phosphoand Evolving Residue-Specific Protein Pho

in Gram-Positive Bacteria

Jakob Fuhrmann,<sup>1,4</sup> Beata Mierzwa,<sup>2</sup> Débora B. Trentini,<sup>1</sup> Silvia Spiess,<sup>3</sup> Anita Lehner,<sup>1</sup> E and Tim Clausen<sup>1,4</sup> "Research Institute of Molecular Pathology (IMP), A-1030 Vienna, Austria <sup>2</sup>Institute of Molecular Biotechnology (IMBA), A-1030 Vienna, Austria <sup>2</sup>Institute of Molecular Biotechnology (IMBA), A-1030 Vienna, Austria <sup>2</sup>The Laboratory for Molecular Infection Medicine Sweden Umeda University, 90187 Umeå, Sweden <sup>3</sup>Present address: Department of Chemistry, The Scripps Hesearch Institute, Jupiter, FL 33458, USA <sup>3</sup>Correspondence: clausen@imp.univie.ac.at. http://dx.doi.org/10.1016/j.celre; 2013 15.023

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of specific client proteins. Bas and function, protein phospl main classes. Phosphatases main classes. Prospiratases nine (pSer, pThr) comprise the tase) and PPM (Mg²-Mn²--t families, whereas enzymes at constitute the protein tyrosine (Barford et al., 1998; Stoker, 2

BACKGROUND: Technologies for making and manipulating DNA have enabled advances in biology ever since the discov-ery of the DNA double helix. But introducing site-specific modifications in the genomes of cells and organisms remained elusive. Early approaches relied on the principle of site-specific recognition of DNA sequences by oligonucleotides, small molecules, or self-splicing introns. More recently, the site-directed zinc finger nucleases (ZFNs) and TAL effector nucleases

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SCIENCE sciencemag.org

widespread adoption of these engineered nucleases for routine use.

ADVANCES: The field of biology is now experiencing a transformative phase with the advent of facile genome engineering in animals and plants using RNA-programmable CRISPR-Cas9. The CRISPR-Cas9 technology originates from type II CRISPR-Cas systems, which provide bacteria with adaptive immunity to viruses and plasmids. The CRISPRassociated protein Cas9 is an endonuclease

peptide nucleic acids (PNAs) and polyamides, were shown to enable targeted binding of chromosomal loci that could be modified if the chemical recognition agent was coupled to a cleavage reagent such as bleomycin (18–20) Another strategy that relied on nucleic acid base pairing was the use of self-splicing introns to change sequences at the DNA (21, 22) or RNA (23) level. Although these approaches did not lead to robust methods, they demonstrated the utility of base pairing for site-specific genome modification.

Corrected 25 November 2014; see full text. 28 N

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## A SIMPLE IDEA



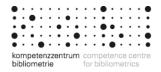


#### A SCALABLE IDEA: THE SCOPUS DATABASE





Data accessed via:





Source: https://www.elsevier.com/\_\_data/assets/pdf\_file/0007/69451/Scopus\_ContentCoverage\_Guide\_WEB.pdf

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#### **OUTLINE**

- BACKGROUND ON THE DATA
- MIGRATION TRENDS AND PATTERNS
- GENDER INEQUALITIES AND INTERNATIONAL MOBILITY
- POLICY SHOCKS AND RETURN MIGRATION



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#### **DATA QUALITY**



- Author name disambiguation in Scopus:
  - 98.3% of author profiles do not include publications written by someone else;
  - 90.6% of author profiles include all publications written by the author.
- Organization disambiguation:

Research Organization Registry (ROR) API

Subset of data with the highest quality:

Period:1996-2020; Type of publications: Articles and Reviews;

→ 36+ million publications



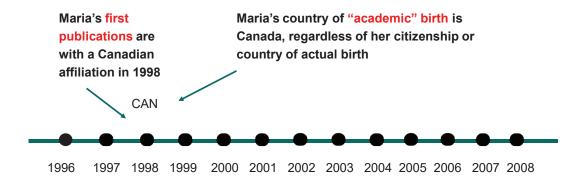
### **IDENTIFYING MIGRATION EVENTS: ILLUSTRATIVE EXAMPLES**



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## **IDENTIFYING MIGRATION EVENTS: ILLUSTRATIVE EXAMPLES**



The country of residence is inferred as the modal country of publications in a given year



### **IDENTIFYING MIGRATION EVENTS: ILLUSTRATIVE EXAMPLES**

Maria changes residence from Canada to the US between 2001 and 2002

CAN CAN CAN USA USA USA USA USA USA USA

1996 1997 1998 1999 2000 2001 2002 2003 2004 2005 2006 2007 2008

Maria's modal countries of publications over time

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### **IDENTIFYING MIGRATION EVENTS: ILLUSTRATIVE EXAMPLES**

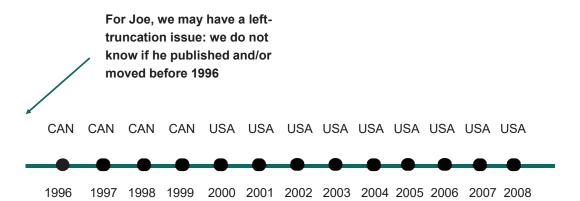
Now, let's consider Joe

CAN CAN CAN USA USA USA USA USA USA USA USA USA

1996 1997 1998 1999 2000 2001 2002 2003 2004 2005 2006 2007 2008



## **IDENTIFYING MIGRATION EVENTS: ILLUSTRATIVE EXAMPLES**



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## **ESTIMATING GENDER USING FIRST NAMES**



#### **ESTIMATING GENDER USING FIRST NAMES**

- Core Dictionary: World Gender-Name Dictionary (WGND), which includes 6.2 million names from 182 countries
- If the name is still missing check other tools like genderize.io
- Validate the results against manually curated dictionaries for 30k+ names

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## **VALIDATION AGAINST MANUALLY CURATED DATA SETS**

Of the names that we classified as female, 90% are actually female

		Indian names	Arabic /Persian names	Japanese names	Chinese names	German names	Russian names
Precision	female	90.49%	94.91%	90.49%	50.55%	98.35%	97.73%
	male	80.92%	95.39%	91.26%	87.60%	97.54%	96.83%
Recall	female	96.25%	96.96%	92.34%	50.30%	98.06%	99.23%
	male	96.22%	92.89%	98.72%	55.05%	98.00%	97.75%

Out of all female names, we correctly classified 96% as female



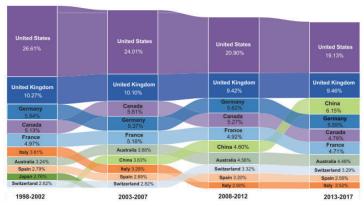
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#### **TOP 10 DESTINATION COUNTRIES**





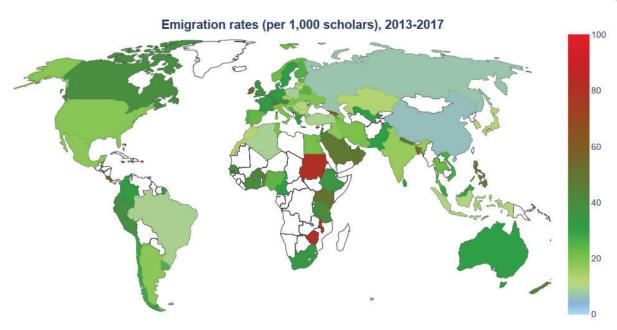
United States
25.93%
United States
20.56%
United Kingdom
8.96%
United Kingdom
8.96%
United Kingdom
8.05%
Cernany
7.03%
Cernany
6.49%
China
1.34%
China
1.34%
China
1.35%
Canada
4.05%
4.05%
Canada

Male migrant researchers

Female migrant researchers





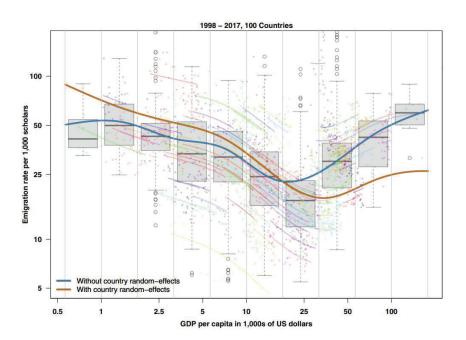


Sanliturk, Zagheni, Danko, Theile, Akbaritabar (2023) PNAS

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## MIGRATION AND ECONOMIC DEVELOPMENT



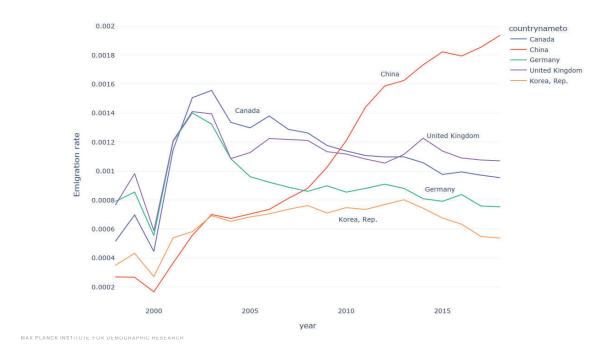


## INTERNATIONAL OUT-MIGRATION FROM THE UNITED STATES





Migration flows from United States to the top 5 destination countries

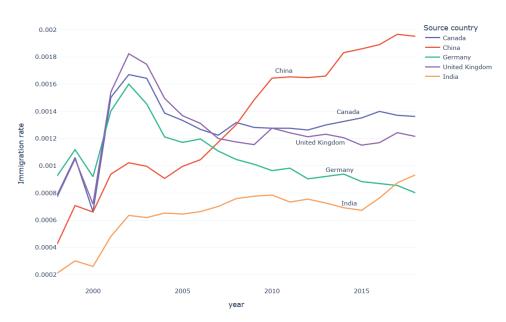


### INTERNATIONAL IN-MIGRATION TO THE UNITED STATES





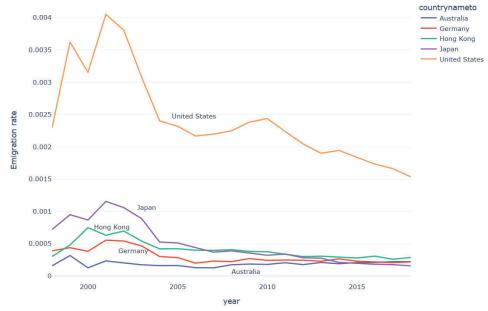
Migration flows from the top 5 source countries to the United States



### INTERNATIONAL OUT-MIGRATION FROM CHINA



Migration flows from China to the top 5 destination countries

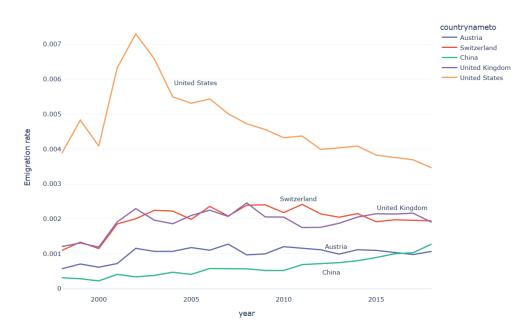


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## INTERNATIONAL OUT-MIGRATION FROM GERMANY



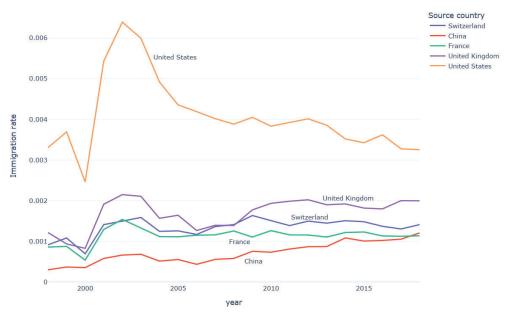
Migration flows from Germany to the top 5 destination countries



## INTERNATIONAL IN-MIGRATION TO GERMANY



Migration flows from the top 5 source countries to Germany



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## **NET MIGRATION RATE - GERMANY**

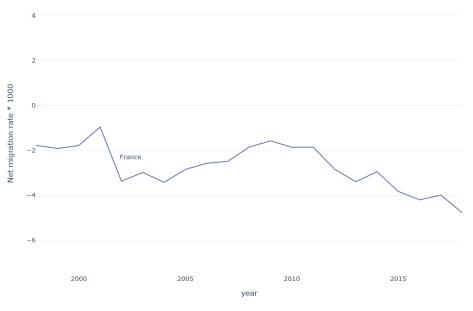
Net migration rates for Germany



## **NET MIGRATION RATE - FRANCE**



Net migration rates for France



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## **NET MIGRATION RATE - SPAIN**

Net migration rates for Spain



## **NET MIGRATION RATE - ITALY**



Net migration rates for Italy

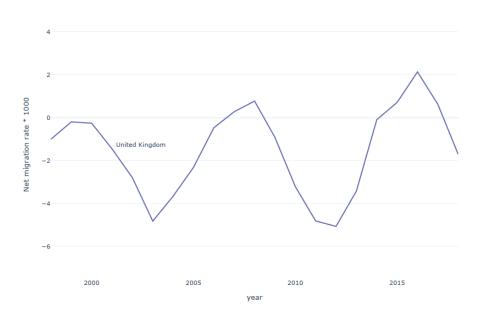


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## **NET MIGRATION RATE - GREAT BRITAIN**



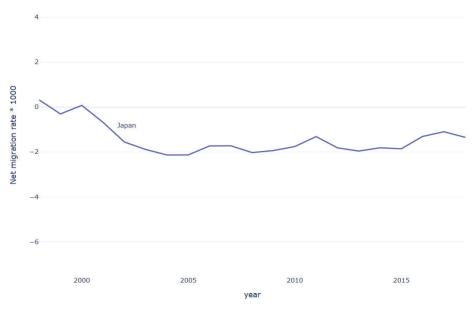
Net migration rates for United Kingdom



## **NET MIGRATION RATE - JAPAN**



Net migration rates for Japan

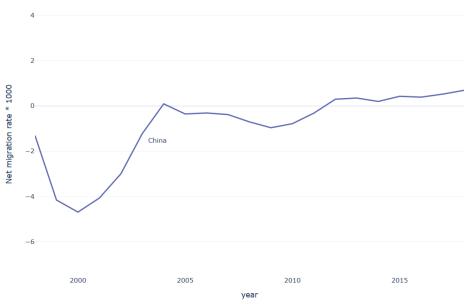


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## **NET MIGRATION RATE - CHINA**



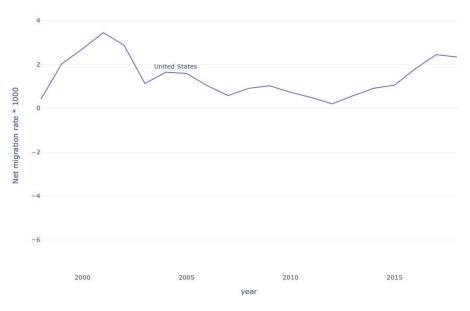
Net migration rates for China



## **NET MIGRATION RATE - USA**



Net migration rates for United States



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## **NET MIGRATION RATE - GERMANY**



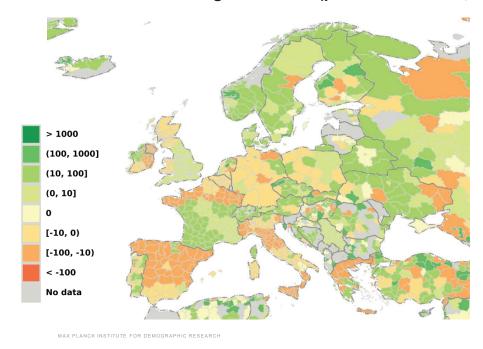
Net migration rates for Germany



#### HETEROGENEITY AT THE SUBNATIONAL LEVEL



#### **Subnational net migration rates (per 1000 scholars, 2012-2017)**



#### Most attractive

FR: Nouvelle-Aquitaine, Occitanie,

Burgogne

PL: Podkarpackie, Kujawsko-

Pomorskie, Lubuskie ES: La Rioja, Navarra

IT: parts of Trentino-South-Tyrol,

Lombardy, Molise DE: Brandenburg

## Losing scholars

FR: Northern

ES: Almost whole country DE: Thuringia, Mecklenburg-

Vorpommern

IT: Most of the country

Akbaritabar et al.(2023) MPIDR Working Paper

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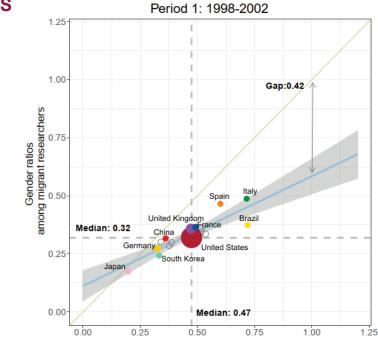
## **WOMEN'S REPRESENTATION IN INTERNATIONAL MOBILITY**

- Progress has been made towards gender equality in science, but women continue to face considerable barriers to participating and advancing in the academic labor force
- International mobility has been recognized as a strategy for scientists to expand their networks and visibility and advance professionally
- International mobility could help narrow the gender gap in academic careers (or could amplify it)

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## WOMEN TO MEN RATIOS FOR SCHOLARS AND MIGRANT SCHOLARS Period 1: 1998-2002

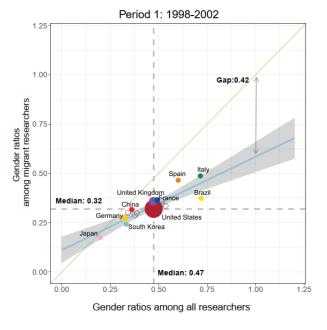


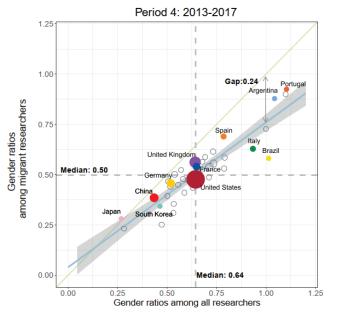


Gender ratios among all researchers

## GENDER RATIOS FOR MIGRANTS CONVERGING TO THOSE OF THE GENERAL POPULATION OF SCHOLARS







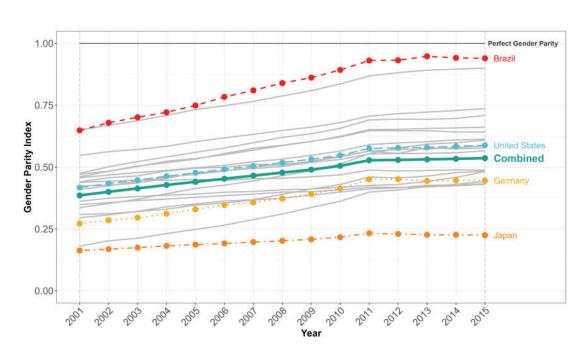
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Zhao, Akbaritabar, Kashyap, Zagheni (2023) PNAS

#### TRENDS IN GENDER PARITY









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#### FLOWS FROM GERMANY AND RETURN RATES



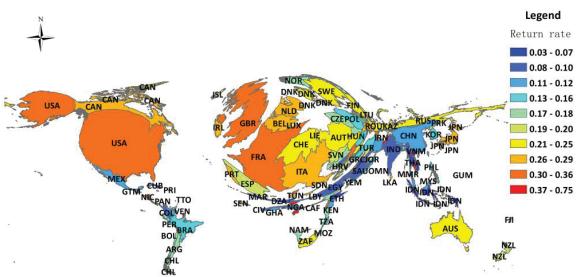
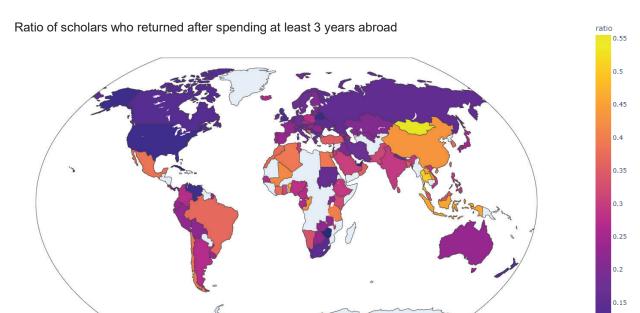


Fig. 2 Outward flows (from Germany) and respective return rates across countries. The sizes of the countries are proportional to the flows of outward researchers from Germany. The colors indicate the differences in the return rates of the German-affiliated researchers returning to Germany from each country.

### **RETURN MIGRATION - COHORT OF 2000**

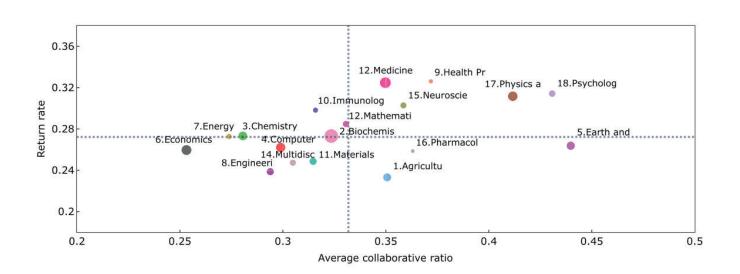




## RETURN RATE TO GERMANY AND FRACTION OF PUBLICATIONS WITH SCHOLARS BASED AT GERMAN INSTITUTIONS







## POLICY SHOCKS: HOW DID BREXIT AFFECT MIGRATION OF SCHOLARS?

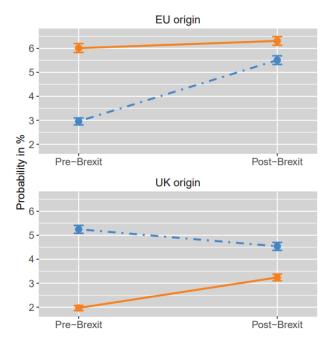




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## POLICY SHOCKS: HOW DID BREXIT AFFECT MIGRATION OF SCHOLARS?





Entering the UK - Leaving the UK



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#### **SOME REFLECTIONS**



- Germany is among the top four worldwide destination countries for scientists and has benefited from playing a central role in the mobility of scholars across Europe and North America.
- · Valuable investments in research are paying off and should continue.
- Key opportunities for the future include strengthening exchange with the Global South and its pool of talent
- Further increasing the representation of women in the global circulation of talent should also be a goal as Germany has room for improvement in this dimension.
- China has been changing the landscape of scientific mobility. The future may depend also on broader geopolitical constellations
- · Not one Germany, but diversity of contexts and heterogeneity across regions
- Policy shocks like Brexit may create disruption. Ultimately science is at its best when circulation and recombination of ideas is favored



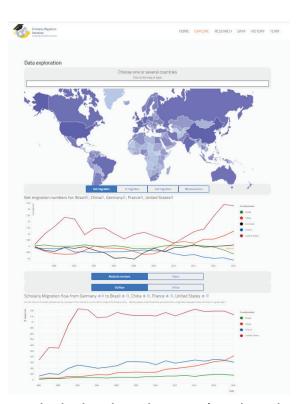


## **Scholarly Migration Database**

Quantifying the Mobility of Scholars

www.scholarlymigration.org









www.scholarlymigration.org/exploration.html

### THANK YOU!

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www.demogr.mpg.de



