

# LUCIA



Lung Cancer-related risk factors and their Impact Assessment

HORIZON-MISS-2021-CANCER-02

## LUCIA Workshop – Understanding Lung Cancer

San Sebastian, Sept. 5<sup>th</sup>, 2023

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**PI: Prof. Dr. Hermann Brenner**  
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# Epidemiology and Lung Cancer Risk Factors

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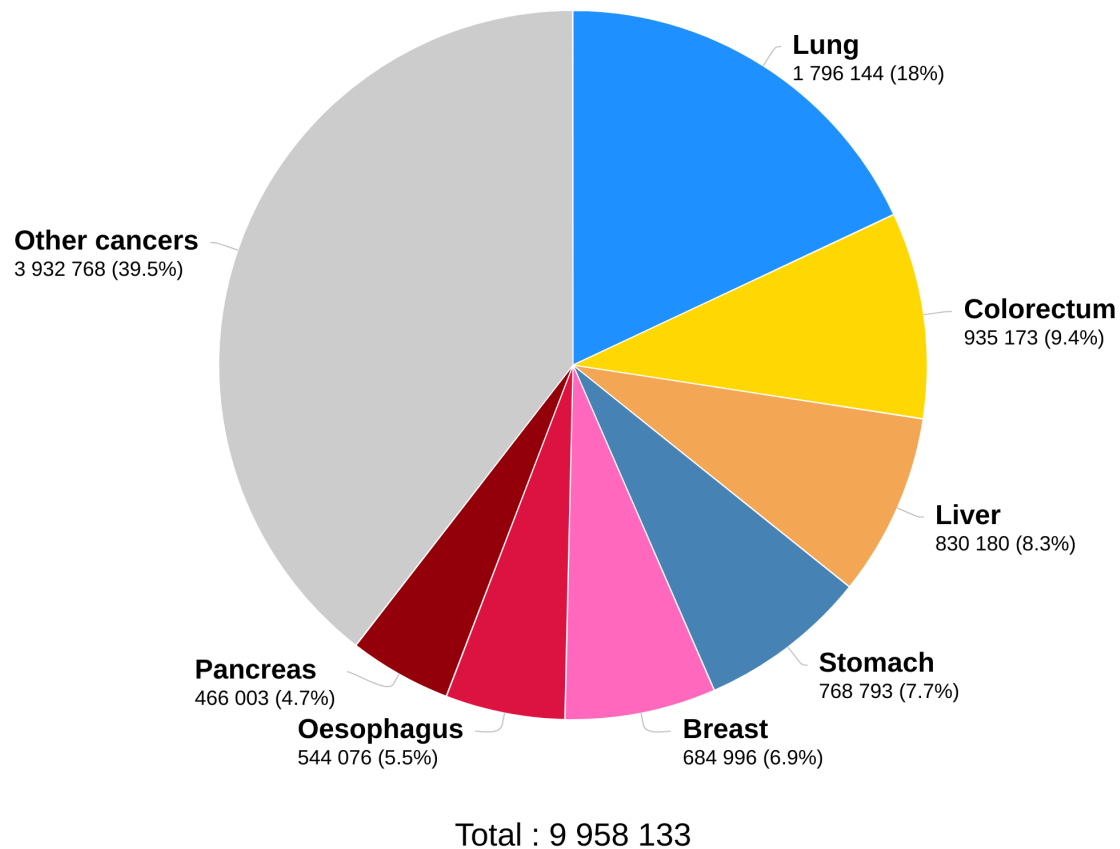


Research for a Life without Cancer

## Outline:

- Epidemiology of lung cancer
- Risk factors: smoking and environmental / occupational exposures
- Regional variability
- Risk-based screening:
  - Criteria-based LDCT-based screening → risk prediction models
  - Blood-based biomarkers

# Estimated number of deaths in 2020, World, both sexes, all ages



- Stage of diagnosis heavily influences treatment outcomes
  - Five-year survival decreases from 55% to 4% when diagnosed in stage IV (versus stage I)

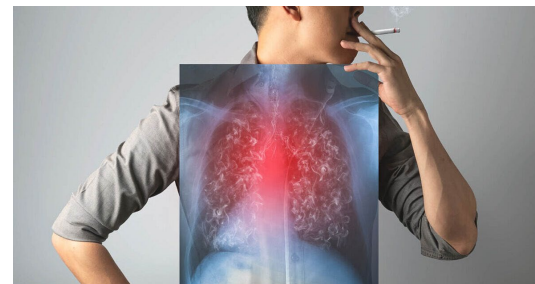
(Cronin et al, *Cancer*. 2018; 124(13):2785-2800)



- Opportunity to reduce mortality by improving primary and secondary prevention (i.e. risk factor reduction, screening and early diagnosis)

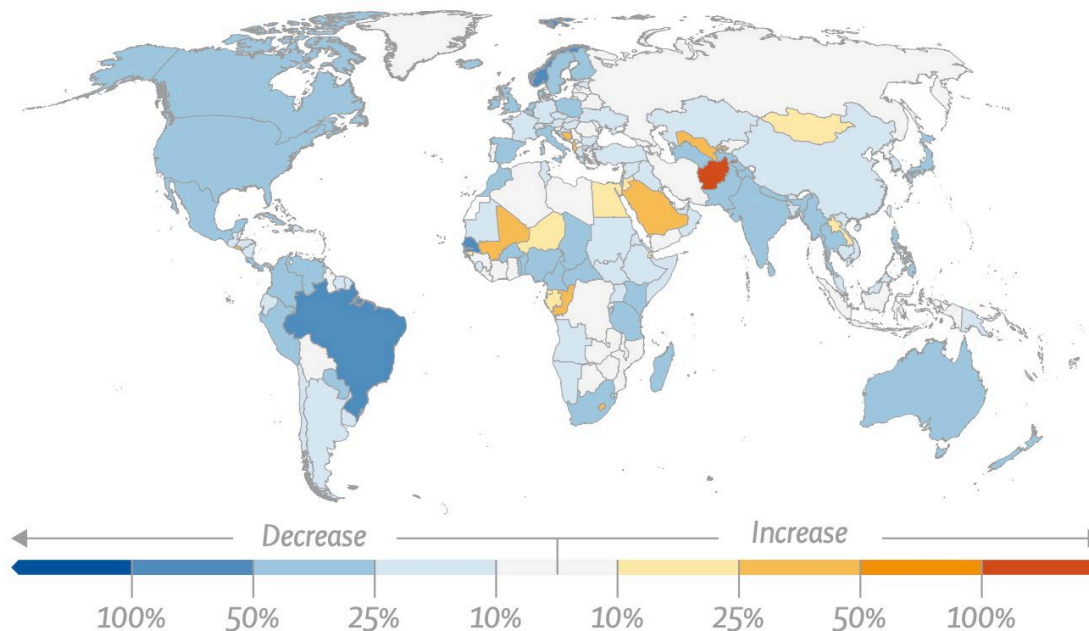
## Tobacco smoking

- **80-90%** of lung cancer cases attributable to smoking
- Increases risk by **15 to 30** times
- **Intensity, duration** and **time since cessation** impact risk
- About **20%** of smokers develop lung cancer
- Second-hand smoke increases risk by **20-30%**





## Change in prevalence by country, 1990–2019



Retrieved from: [https://www.thelancet.com/pb/assets/raw/Lancet/infographics/tobacco/Tobacco\\_infographics\\_global.pdf](https://www.thelancet.com/pb/assets/raw/Lancet/infographics/tobacco/Tobacco_infographics_global.pdf)

## 🔍 Globally, prevalence of tobacco smoking is estimated at **22%**

- Regions with the highest prevalence: South-East Asia, Western Pacific, Europe (estimated prevalence of 25%-30%)
- In Europe:
  - Men (estimated prevalence of 33%), Women (estimated prevalence of 18%)
  - Slower decline in cigarette smoking in women than in men

## 📈 E-cigarettes use has increased among adolescents and young adults



WHO global report tobacco, 2019



## Other risk factors for lung cancer

### **Environmental and occupational exposures:**

- Unprocessed biomass fuels
- Arsenic
- Radon
- Asbestos
- Diesel exhaust

### **Family history and other diseases:**

- Genetic variants
- Previous chronic lung diseases
- HIV / HPV infection

### **Lifestyle factors:**

- Diet and metabolic factors

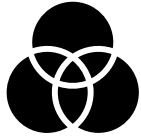


## Incidence and mortality trends vary according to:

- National human development index (HDI):
  - Mortality rate in very high HDI (22.8 ASR per 100,000 py) versus low HDI settings (3.2 ASR per 100,000 py)
- World region:
  - Eastern Asia, Western Europe and Northern America had among the highest incidence rates (32.6-34.4 ASR per 100,000 py)
- Sex:
  - Trends in incidence and mortality were decreasing among men in most countries
  - In European countries, trends increased in women

- Regional variation in lung cancer incidence and mortality linked to

- **Differences** in smoking patterns



- **Environmental** exposures

- Frequency of somatic **mutations** (such as (EGFR) alterations)

- Lung cancer **subtypes**



- Existing predictive models for risk stratification:

Supplementary Table 1: Covariates included in the established LC risk prediction models

	Model [reference]								
	Bach Model [14]	Spitz Model [15]	LLP Model [16]	Hoggart Model [17]	PLCO <sub>M</sub> 2012 [18]/ PLCO <sub>A</sub> 2014 [19] Models	Pittsburgh Predictor[21]	LLP <sub>i</sub> [20]/ LLP <sub>v3</sub> [23] Models	LCRAT [22]/ LCDRAT[22]	
Country/Region	USA	USA	UK	EU	USA	USA	UK	USA	
Target population, smoking status	Ever smokers	Never, former & current smokers	Individuals aged	Ever smokers	Ever Smokers/ General population	Ever smokers	Individuals aged	Ever smokers	
Target population, age range (years)	44-75	≥ 18	20-80	40-65	55-74/ 55-80	55-74	45-79/ 50-79	55-74/ 55-74	
Cumulative risk prediction (year/s)	5	1	5	1	6/ 6	6	8.7/ 5	5/ 5	
Covariates included	<b>Demographics</b>								
Age	X	X	X	X	X	X	X	X	X
Gender	X	X	X				X	X	X
Race/ethnicity <sup>a</sup>					X				X
BMI					X				X
Education					X				X
Smoking									
Smoking status		X		X	X	X			
Years/age quit smoking	X	X		X	X				X
Years smoked	X		X	X	X	X	X	X	X
Cigarettes per day	X			X	X	X	X	X	X
Pack-years		X							X
Comorbidities									
Lung disease (COPD/Emphysema)		X			X		X	X	X
Prior cancer			X		X		X		
Pneumonia			X						
Hayfever		X							
Occupational exposures									
Asbestos exposure	X	X	X						
Dust exposure		X							
Family history									
Any FDR with cancer		X							
Any FDR with smoking related cancer		X							
Any FDR with LC			X		X		X	X	X
Num. FDR with LC									X
Early onset FDR with LC			X				X		

Bhardwaj et al  
Lung Cancer 2022, 174:83-90

## ESTHER Study cohort (9,940 participants)

Saarland, Germany:

N = 4812 Ever smokers aged 50-75 years  
(with no previous LC diagnoses)

17 year FU

262 Incident LC, 4450 No LC

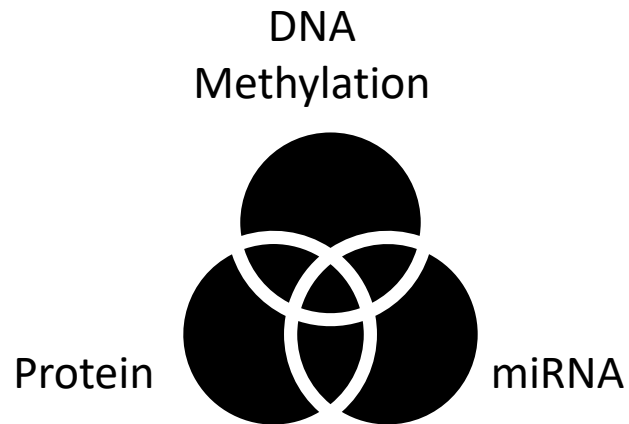
Bhardwaj et al  
Lung Cancer 2022, 174:83-90

**Table 2**

Performances of different **LC risk prediction models** for incident cases identified during 6, 11 and 17 years of follow-ups among **ever smoking participants** of the ESTHER-study.

Model, country [reference]	Time horizon (n total = 4812)		
	0-6 years (n iLC = 102)	0-11 years (n iLC = 179)	0-17 years (n iLC = 262)
	AUC (95% CI)		
Bach model, USA [14]	0.782 (0.739-0.824)	0.774 (0.742-0.805)	0.766 (0.740-0.792)
Spitz model, USA [15]	0.725 (0.677-0.774)	0.688 (0.648-0.728)	0.638 (0.604-0.672)
LLP model, UK [16]	0.744 (0.698-0.790)	0.717 (0.679-0.756)	0.676 (0.641-0.710)
Hoggart model, Europe [17]	0.705 (0.649-0.761)	0.708 (0.667-0.749)	0.712 (0.678-0.745)
PLCO <sub>M2012</sub> , USA [18]	0.751 (0.699-0.803)	0.742 (0.703-0.781)	0.739 (0.707-0.771)
PLCO <sub>all2014</sub> model, USA [19]	0.749 (0.698-0.800)	0.741 (0.702-0.779)	0.742 (0.711-0.774)
LLPi model, UK [20]	0.773 (0.731-0.815)	0.753 (0.720-0.787)	0.725 (0.697-0.754)
Pittsburgh Predictor, USA [21]	0.778 (0.737-0.820)	0.763 (0.730-0.797)	0.766 (0.739-0.792)
LCRAT, USA [22]	0.786 (0.745-0.827)	0.771 (0.739-0.804)	0.771 (0.745-0.797)
LCDRAT, USA [22]	0.787 (0.746-0.828)	0.770 (0.738-0.802)	0.765 (0.739-0.792)
LLPv3 model, UK [23]	0.773 (0.731-0.815)	0.753 (0.720-0.787)	0.725 (0.697-0.754)

## Improvements of risk models with blood-based biomarkers



## miRNA-Score → increase in AUCs of risk models of 0.041 and 0.096

**TABLE 1** Performance of LC models\* [7] individually and in combination with the miR-score for LC risk prediction among ever-smokers ( $n_{\text{cases/controls}} = 206/101$ ) in prospective set participants

LC model	AUC (95% CI)			P-value for improvement in AUC
	LC model only	LC model + miR-score apparent <sup>#</sup>	LC model + miR-score .632+ <sup>\$</sup>	
LLPi	0.712 (0.651-0.774)	0.765 (0.708-0.821)	0.762	0.019
Pittsburgh Predictor	0.717 (0.656-0.779)	0.764 (0.707-0.821)	0.762	0.042
Bach	0.716 (0.654-0.778)	0.760 (0.703-0.817)	0.758	0.034
PLCO <sub>M2012</sub>	0.694 (0.632-0.755)	0.757 (0.700-0.814)	0.754	0.008
LLP	0.669 (0.606-0.732)	0.750 (0.693-0.808)	0.747	0.003
Hoggart	0.655 (0.591-0.719)	0.738 (0.679-0.797)	0.734	0.004
Spitz	0.623 (0.557-0.689)	0.723 (0.665-0.781)	0.719	0.001
LCRAT	0.721 (0.658-0.783)	0.765 (0.707-0.822)	0.762	0.061

Yu et al Cancer Communications 2022, 42(11), 1222–1225

Inflammatory plasma protein biomarker score → increase in AUCs of risk models up to 0.07

Ever smoking participants from ESTHER study, incl. 172 incident LC cases and 285 randomly selected participants free of LC

92 inflammation protein biomarkers were measured

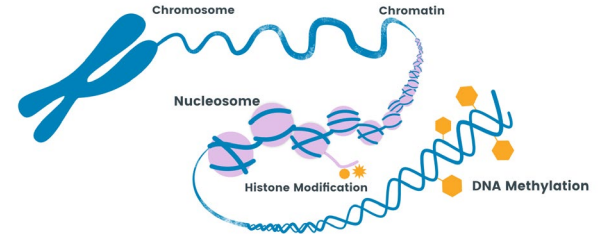
0.632+ bootstrap applied for correction of overoptimism



## DNA Methylation analyses in ESTHER Dataset

N = 162 ever-smoking cases,

N = 721 ever-smoking randomly selected controls



### AHRR (cg05575921) methylation

- Adding to LC risk models, increase in AUCs from 0.036 to 0.133 (P < 0.05 for all 10 risk models)

### F2RL3 (cg03636183) methylation

- Adding to LC risk models, increase in AUCs from 0.041 to 0.129 (P < 0.05 for all 10 risk models)

## Future epidemiological work within LUCIA

- Analyses in ESTHER dataset
  - Complete genotyping
  - Methylation arrays for all lung cancer cases and controls
  - microRNA analyses
  - OLINK proteomics analyses



## Conclusions:

- Globally lung cancer is the leading cause of cancer deaths, with geographical variability in incidence and mortality
- Regional variability is largely reflected by trends in tobacco smoking and other risk factors
- Risk models for selecting high-risk individuals for screening are promising, especially in combination with biological signatures



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## Evidence from LDCT-based screening program trials:

### US National Lung Screening Trial (NLST) (US) and NELSON (EU)

- Enrollment criteria-based (age, history of smoking, and time since smoking cessation)
- 20% reduction in mortality but still substantial number of false-negative and false-positive results → need for optimized selection for screening
- Slow uptake when compared with other cancer screening programs



## • Lung cancer risk factor tobacco smoking

- Impact of smoking on lung cancer affected by intensity, duration and time since cessation
- Less than 20% of smokers develop lung cancer (IARC, 2004) – many factors determine individual susceptibility

