Lecture 11: AI and COVID-19

Serena Yeung

BIODS 220: AI in Healthcare

Announcements

- Project milestone due Fri Oct 30
- Project milestone presentations Mon Nov 2 in-class
 - 4 minutes per group, strict time limit. It's ok to have a subset of group members present
 - Should summarize all components of milestone report (5 pts total)
 - Pre-recorded video option can be requested for those unable to attend
 - See Piazza post for more details about all of this

Today

 Applications of AI in Healthcare through the lens of a real-world case study: COVID-19



First application area: AI interpretation of chest radiology images



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Bernheim et al.: COVID-19 hallmarks on chest CTs based on radiologist review

Key Results

- The hallmark CT manifestations of coronavirus disease 2019 (COVID-19) include bilateral and peripheral ground-glass and consolidative pulmonary opacities, sometimes with a rounded morphology and peripheral lung distribution.
- As the time between onset of symptoms and initial chest CT increases, some CT findings are observed with increasing frequency, including consolidation, bilateral and peripheral lung disease, greater total lung involvement, linear opacities, and the appearance of a crazy-paving pattern and reverse halo sign.
- Certain chest CT findings, including pleural effusions, lymphadenopathy, pulmonary nodules, and lung cavitation, are characteristically absent, and more than half of patients imaged quickly after symptom onset have a normal CT scan.

Bernheim et al. Chest CT Findings in Coronavirus Disease 2019 (COVID-19): Relationship to Duration of Infection, 2020.

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Bernheim et al.: COVID-19 hallmarks on chest CTs based on radiologist review

Bernheim et al. Chest CT Findings in Coronavirus Disease 2019 (COVID-19): Relationship to Duration of Infection, 2020.



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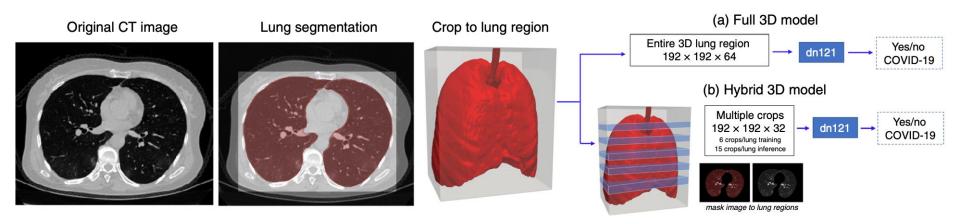
Figure 1: (a) Axial CT image obtained without intravenous contrast material in a 36-year-old man shows bilateral ground-glass opacities in upper lobes with a rounded morphology (arrows). (b) Axial CT image obtained in a 44-year-old man shows larger ground-glass opacities in the bilateral lower lobes with a rounded morphology (arrows). (c) Axial CT image obtained in a 65-year-old woman shows bilateral ground-glass and consolidative opacities with a striking peripheral distribution (arrows).

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- Detection of COVID-19 from CT images
- 2 stage process: lung segmentation followed by classification of COVID-19 or not
- Multinational dataset of 2724 scans from 2617 patients, with 1029 scans (922) patients confirmed positive for COVID-19

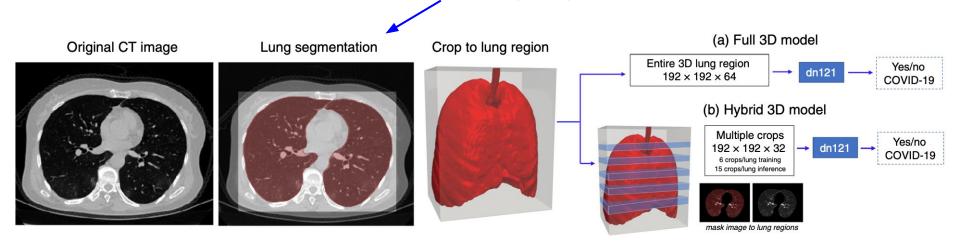


Harmon et al. Artificial intelligence for the detection of COVID-19 pneumonia on chest CT using multinational datasets, 2020.

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- Detection of COVID-19 from CT images
- 2 stage process: lung segmentation followed by classification of COVID-19 or not
- Multinational dataset of 2724 scans from 2617 patients, with 1029 scans (922) patients confirmed positive for COVID-19
 First stage: segmentation



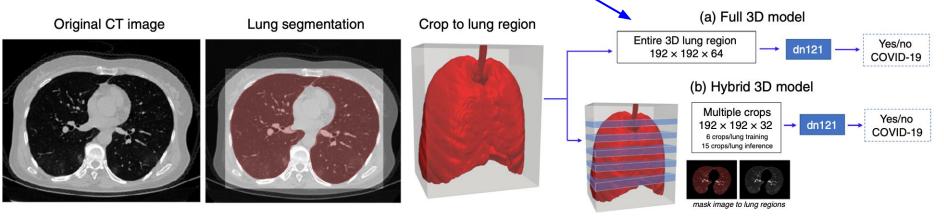
Harmon et al. Artificial intelligence for the detection of COVID-19 pneumonia on chest CT using multinational datasets, 2020.

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- Detection of COVID-19 from CT images
- 2 stage process: lung segmentation followed by classification of COVID-19 or not
- Multinational dataset of 2724 scans from 2617 patients, with 1029 scans (922) patients confirmed positive for COVID-19
 Second stage: classification based on whole lung

region vs. combination of cropped regions

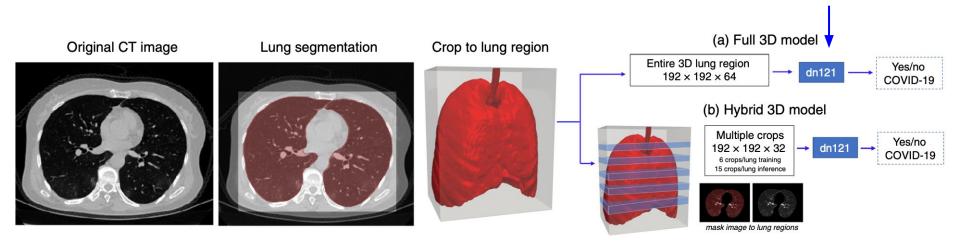


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- Detection of COVID-19 from CT images
- 2 stage process: lung segmentation followed by classification of COVID-19 or not
- Multinational dataset of 2724 scans from 2617 patients, with 1029 scans (922) patients confirmed positive for COVID-19
 "DenseNet" convolutional neural network architecture



Harmon et al. Artificial intelligence for the detection of COVID-19 pneumonia on chest CT using multinational datasets, 2020.

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Multinational patient dataset

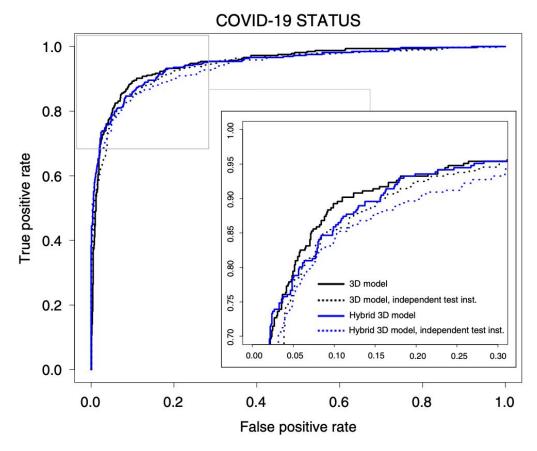
Disease cohort	Center	Demographics	Training	Validation	Testing
COVID-19	Hubei, China	363 Male, 353 female	369 Scans	122 Scans	207 Scans
		Median 49 (18ª-92)	354 Patients	113 Patients	207 Patients
	Milan, Italy	220 Male, 90 female	57 Scans	24 Scans	54 Scans
		Median 60 (18-96)	52 Patients	17 Patients	54 Patients
	Tokyo, Japan	91 Male, 60 female	100 Scans	31 Scans	49 Scans
		Median 60 (4-87)	45 Patients	15 Patients	49 Patients
	Milan, Italy	10 Male, 5 female	-	-	15 Scans
		Median 55 (31-85)			15 Patients
	Syracuse, NY, USA	^b See footnote	-	-	1 Scan
					1 Patient
Any clinical indication	Syracuse, NY, USA	437 Male, 534 female	356 Scans	93 Scans	500 Scans
		Median 65 (19-100)	356 Patients	93 Patients	500 Patients
Cancer diagnosis and/or staging	LIDC ²³	N/A	149 Scans	50 Scans	271 Scans
			149 Patients	50 Patients	271 Patients
	NIH, USA	100 Male	-	-	100 Scans
		Median 69 (30-89)			100 Patients
Pneumonia	Syracuse, NY, USA	73 Male, 42 female	-	-	140 Scans
		Median 66 (13-101)			140 Patients
	NIH, USA	28 Male, 8 female	28 Scans	8 Scans	-
		Median 21 (4-71)	28 Patients	8 Patients	
Total			1059 Scans	328 Scans	1337 Scans
			984 Patients	296 Patients	1337 Patients

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Harmon et al. Artificial intelligence for the detection of COVID-19 pneumonia on chest CT using multinational datasets, 2020.

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Model achieved 90.8% accuracy (84% sensitivity, 93% specificity)

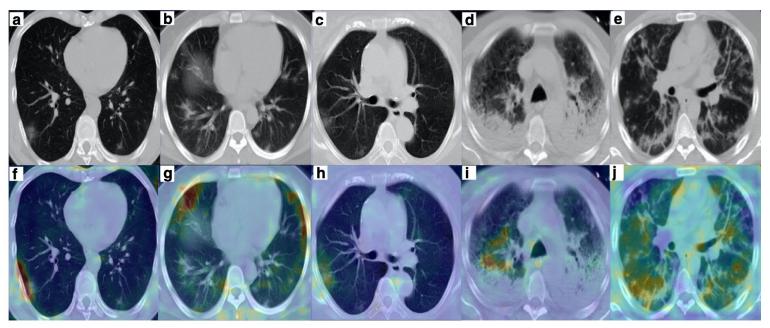


Harmon et al. Artificial intelligence for the detection of COVID-19 pneumonia on chest CT using multinational datasets, 2020.

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Grad-CAM saliency maps showing regions contributing most to model prediction (will discuss more in upcoming lecture)



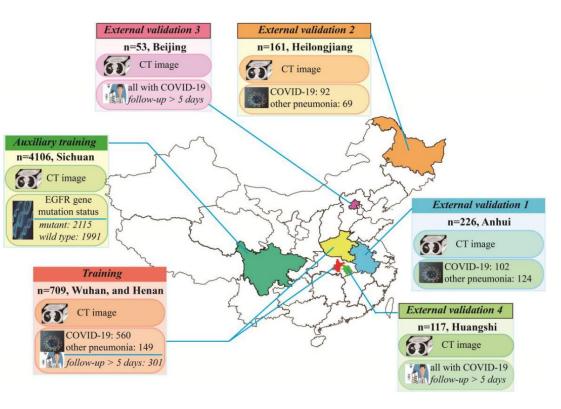
Harmon et al. Artificial intelligence for the detection of COVID-19 pneumonia on chest CT using multinational datasets, 2020.

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Wang S. et al.

- Also detection of COVID-19 from CT images, based on 1266 patients (924 with COVID-19) from 7 Chinese cities or provinces
- Addressed lack of available data due to ongoing pandemic, through pre-training using a different dataset of 4106 lung cancer patients (that was trained to predict accompanying epidermal growth factor receptor (EGFR) gene mutation)



Wang S et al. A Fully Automatic Deep Learning System for COVID-19 Diagnostic and Prognostic Analysis, 2020.

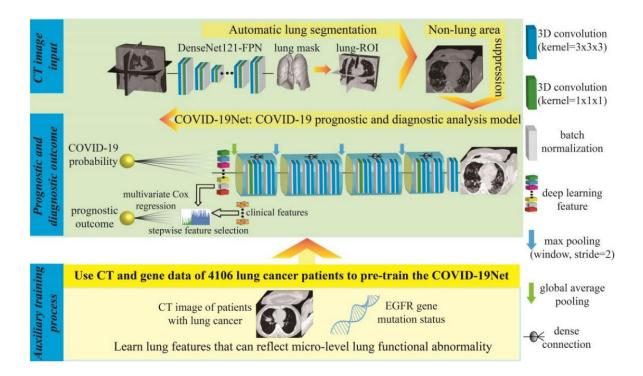
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Wang S. et al.

- Deep learning architecture is similar to a DenseNet (stacked modules, with dense connections between them)

Also extracted
 64-dimensional visual features
 from the classification model,
 and combined it with clinical
 features to train a model for
 prognosis (higher-risk vs.
 lower-risk patients)



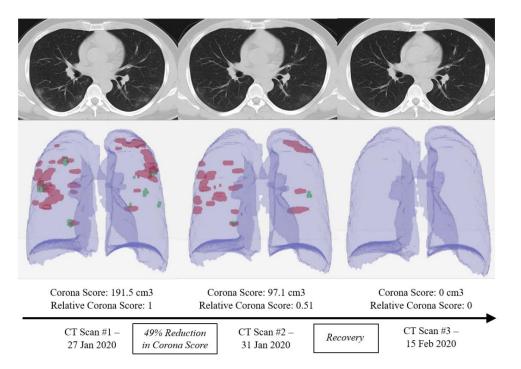
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Wang S et al. A Fully Automatic Deep Learning System for COVID-19 Diagnostic and Prognostic Analysis, 2020.

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Gozes et al.

- Beyond COVID-19 classification on CTs, also outputs a "Corona score" to measure progression of the disease over time
- The score is a volumetric measurement of the opacities burden, and is based on a volumetric summation of network-activation maps and localized nodule detections
- A "relative Corona score" can perform patient-specific monitoring by normalizing the score by the score computed at the first time point.



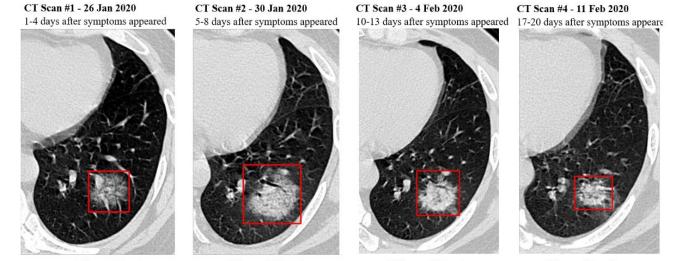
Gozes et al. Rapid AI Development Cycle for the Coronavirus (COVID-19) Pandemic: Initial Results for Automated Detection & Patient Monitoring using Deep Learning CT Image Analysis, 2020.

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Gozes et al.

Multi-time point tracking of patient disease progression



Volume: 9.9 cm3 Average Axial Diameter: 30.3 mm

Volume: 19.9 cm3 Average Axial Diameter: 37.4 mm

Volume: 13.2 cm3 Volume: 6.3 cm3 Average Axial Diameter: 34.2 mm Average Axial Diameter: 26.6 mm

Gozes et al. Rapid AI Development Cycle for the Coronavirus (COVID-19) Pandemic: Initial Results for Automated Detection & Patient Monitoring using Deep Learning CT Image Analysis, 2020.

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Wang L. et al.

- This work takes a different approach and tries to detect COVID-19 from chest x-rays instead of CTs, since x-rays are fast, more accessible (especially in developing countries), and portable (can be performed e.g., within an isolation room)
- Trained a deep learning model to predict no infection, non-COVID-19 infection, COVID-19 infection





No infection

COVID-19 infection

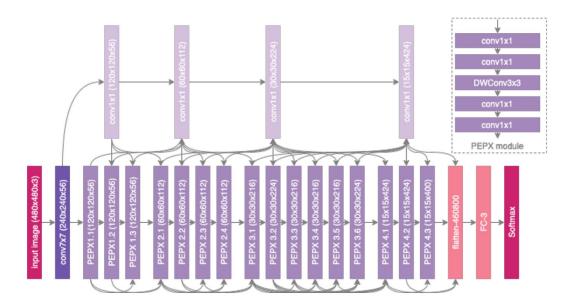
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Wang et al. COVID-Net: A Tailored Deep Convolutional Neural Network Design for Detection of COVID-19 Cases from Chest X-Ray Images, 2020.

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Wang S. et al.

- Model architecture was selected based on a "network generation" approach to design a high-performing network for the task. But still based on familiar components!



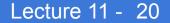
Sensitivity (%)				Positive Predictive Value (%)			
Architecture	Normal	Non-COVID19	COVID-19	Architecture	Normal	Non-COVID19	COVID-19
VGG-19	98.0	90.0	58.7	VGG-19	83.1	75.0	98.4
ResNet-50	97.0	92.0	83.0	ResNet-50	88.2	86.8	98.8
COVID-Net	95.0	94.0	91.0	COVID-Net	90.5	91.3	98.9

Wang et al. COVID-Net: A Tailored Deep Convolutional Neural Network Design for Detection of COVID-19 Cases from Chest X-Ray Images, 2020.

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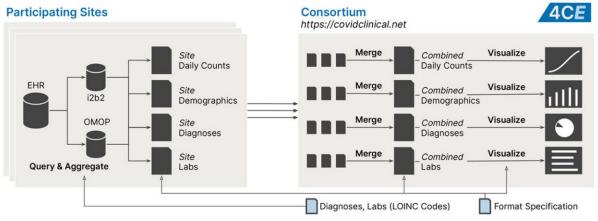
Second application area: Modeling patient outcomes using EHR data





Aggregating data across many hospitals: the 4CE consortium

- Consortium for Clinical Characterization of COVID-19 by EHR (4CE): international consortium of 96 hospitals across five countries
- Used platforms such as OMOP to map all EHR to a common data model
- Total data covers 27,584 COVID-19 cases with 187,802 laboratory tests
- Initially includes 14 laboratory markers of cardiac, renal, hepatic, and immune dysfunction that have been associated with poor outcome in COVID-19 patients



Brat et ak, International electronic health record-derived COVID-19 clinical course profiles: the 4CE consortium, 2020.

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Remember: OMOP Common Data Model

- Observational Medical Outcomes Partnership (OMOP)
- Created from public-private partnership involving FDA, pharmaceutical companies, and healthcare providers
- Standardized format and vocabulary
- Allows conversion of patient data from different sources into a common structure for analysis
- Intended to support data analysis

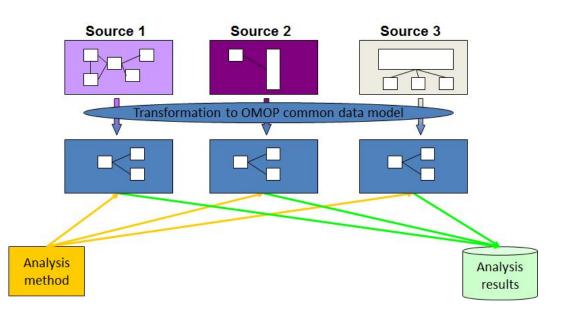


Figure credit: https://www.ohdsi.org/wp-content/uploads/2014/07/Why-CDM.png

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Remember: OMOP Common Data Model

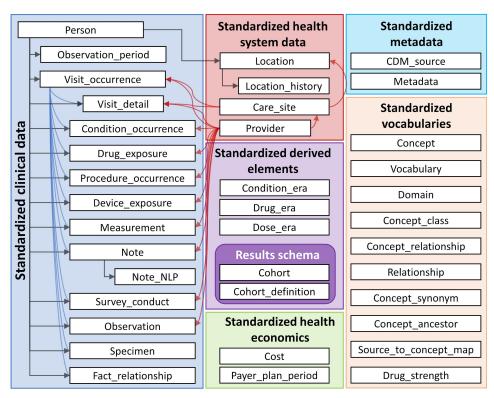


Figure credit: https://ohdsi.github.io/TheBookOfOhdsi/images/CommonDataModel/cdmDiagram.png

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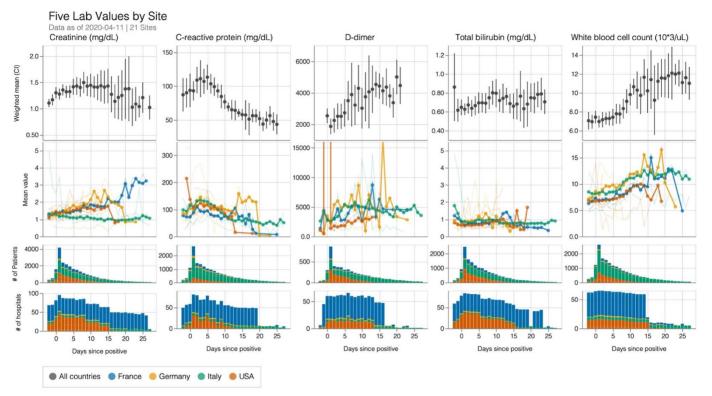
Aggregating data across many hospitals: the 4CE consortium

Healthcare system	Acronym	City	Country	Population	Hospitals	Beds	Inpatient discharges/yea
Assistance Publique—Hôpitaux de Paris	APHP	Paris	France	Adult & Pediatric	39	20,098	1,375,538
Bordeaux University Hospital	FRBDX	Bordeaux	France	Adult & Pediatric	3	2,676	130,033
Erlangen University Hospital	UKER	Erlangen	Germany	Adult & Pediatric	1	1,400	65,000
Medical Center, University of Freiburg	UKFR	Freiburg	Germany	Adult & Pediatric	1	1,660	71,500
University Medicine Mannheim	UMM	Mannheim	Germany	Adult & Pediatric	1	1,352	50,748
ICSM Pavia Hospital	ICSM1	Pavia	Italy	Adult	1	426	8616
ICSM Lumezzane/Brescia Hospitals	ICSM5	Lumezzane/Brescia	Italy	Adult	1	149	1296
ICSM Milano Hospital	ICSM20	Milan	Italy	Adult	1	200	2432
Policlinico di Milano	POLIMI	Milan	Italy	Adult & Pediatric	1	900	40,000
ASST Papa Giovanni XXIII Bergamo	HPG23	Bergamo	Italy	Adult & Pediatric	1	1080	45,000
National University Hospital	NUH	Singapore	Singapore	Adult & Pediatric	1	1556	100,977
Boston Children's Hospital	BCH	Boston, MA	USA	Pediatric	1	404	28,000
Beth Israel Deaconess Medical Center	BIDMC	Boston, MA	USA	Adult	1	673	40,752
Children's Hospital of Philadelphia	CHOP	Philadelphia, PA	USA	Pediatric	1	564	25,406
University of Kansas Medical Center	KUMC	Kansas City, KS	USA	Adult & Pediatric	1	794	54,659
Mayo Clinic	MAYOC	Rochester, MN	USA	Adult & Pediatric	1	2059	100,000
Mass General Brigham (Partners Healthcare)	MGB	Boston, MA	USA	Adult & Pediatric	10	3418	163,521
Medical University of South Carolina	MUSC	Charleston, SC	USA	Adult & Pediatric	8	1600	55,664
University of Pennsylvania	UPenn	Philadelphia, PA	USA	Adult	5	2469	118,188
University of California, LA	UCLA	Los Angeles, CA	USA	Adult & Pediatric	2	786	40,526
University of Michigan	UMICH	Ann Arbor, MI	USA	Adult & Pediatric	3	1000	49,008
University of North Carolina at Chapel Hill	UNC	Chapel Hill, NC	USA	Adult & Pediatric	11	3095	52000
UT Southwestern Medical Center	UTSW	Dallas, TX	USA	Adult	1	608	26,905
				Total	96	45,352	2,444,792

Brat et ak, International electronic health record-derived COVID-19 clinical course profiles: the 4CE consortium, 2020.

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Aggregating data across many hospitals: the 4CE consortium



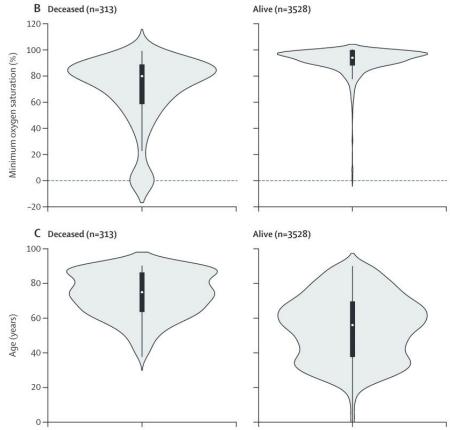
Brat et ak, International electronic health record-derived COVID-19 clinical course profiles: the 4CE consortium, 2020.

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Yadaw et al.

- Prediction of COVID-19 patient mortality from patient clinical variable data
- Trained on 3841 patients from the Mount Sinai Health System in NYC. Tested on 961 retrospective and 249 prospective patients.
- Needed to perform missing value imputation (remember from Lecture 6)



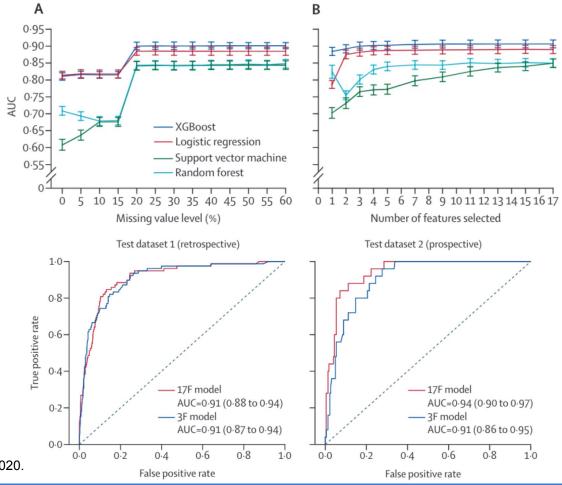
Yadaw et al. Clinical features of COVID-19 mortality: development and validation of a clinical prediction model, 2020.

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Yadaw et al.

Evaluation of multiple types of machine learning models



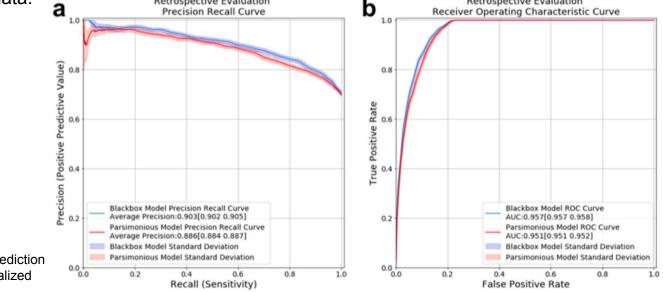
Yadaw et al. Clinical features of COVID-19 mortality: development and validation of a clinical prediction model, 2020.

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Razavian et al.

- Instead of mortality, predict favorable outcomes (may be more meaningful when ICUs are already saturated!)
- Trained multiple types of machine learning models on 3345 and 474 prospective hospitalizations, using clinical variable data. Retrospective Evaluation



Razavian et al. A validated, real-time prediction model for favorable outcomes in hospitalized COVID-19 patients, 2020.

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Razavian et al.

Integrated and deployed in NYU hospitals EHR system

Covid-19 Low Risk Patient Name of Adverse Events 100 Covid-19 Low Risk of Adverse Events within 96 hours Factors Contributing to Score 98 61% O2 Device is greater than Nasal cannula 100 9% RR Max in last 12 hr is 36 7% Blood urea nitrogen Last is 39 21 4% C-reactive protein Last is 235 High 2% HR Min in last 12 hr is 77 19 1% Temp Max in last 12 hr is 98.2 100 -4% Nasal cannula O2 flow rate Max in last 12 hr is N/A 19 -4% SpO2 Min in last 12 hr is 91 -8% Eosinophils % Last is 7 6 6 Covid-19 Low Risk of Adverse Events within 96 hours Factors Contributing to Score 13% Blood urea nitrogen Last is 24 6 8% RR Max in last 12 hr is 20 7% HR Min in last 12 hr is 93 3% RR Min in last 12 hr is 18 Low 3% LDH Last is 713 -8% Platelet count Last is 240 15 -9% O2 Device is None (Room air) -19% Eosinophils % Last is 5 -29% SpO2 Min in last 12 hr is 97 6 (c) Epic Systems Corporation. Used with permission

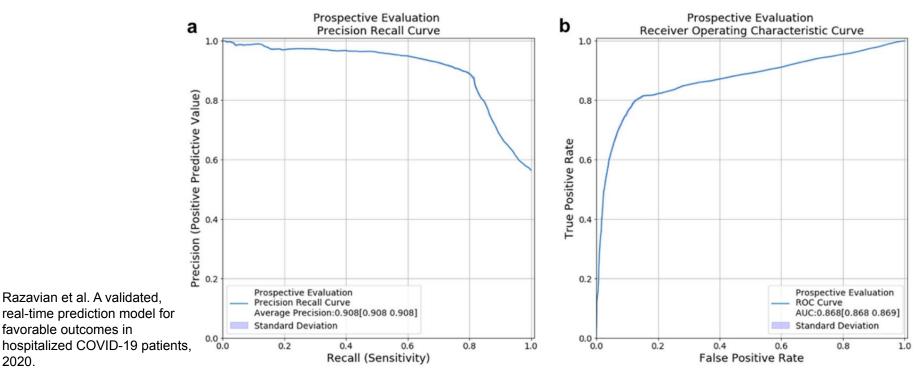
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Razavian et al. A validated, real-time prediction model for favorable outcomes in hospitalized COVID-19 patients, 2020.

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Razavian et al.

Prospective model performance



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2020.

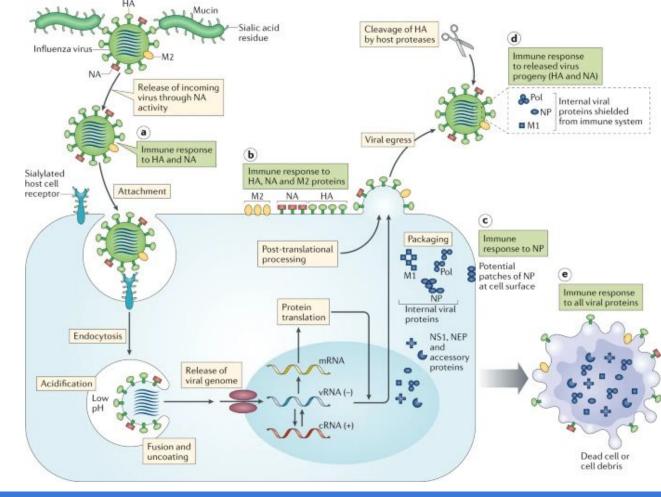
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Third application area: Finding treatments for the disease



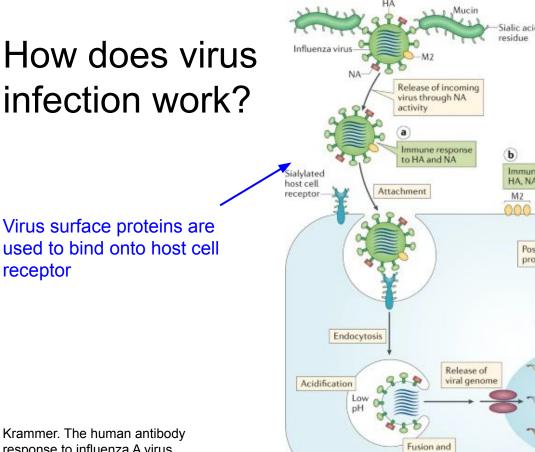
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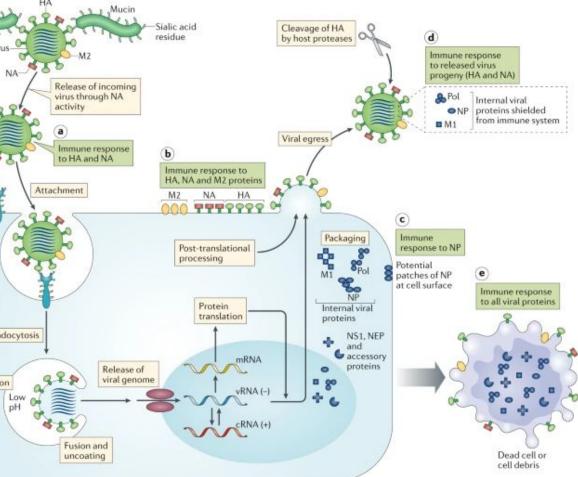
Krammer. The human antibody response to influenza A virus infection and vaccination, 2020.



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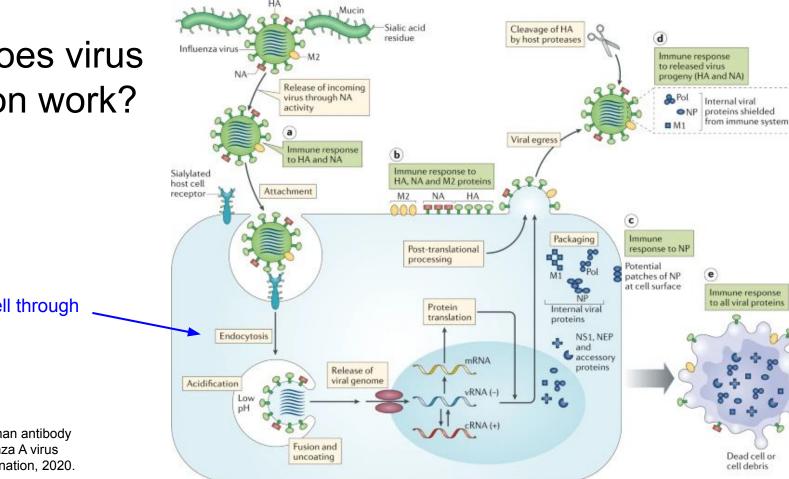
response to influenza A virus infection and vaccination, 2020.

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Virus enters cell through endocytosis

Krammer. The human antibody response to influenza A virus infection and vaccination, 2020.

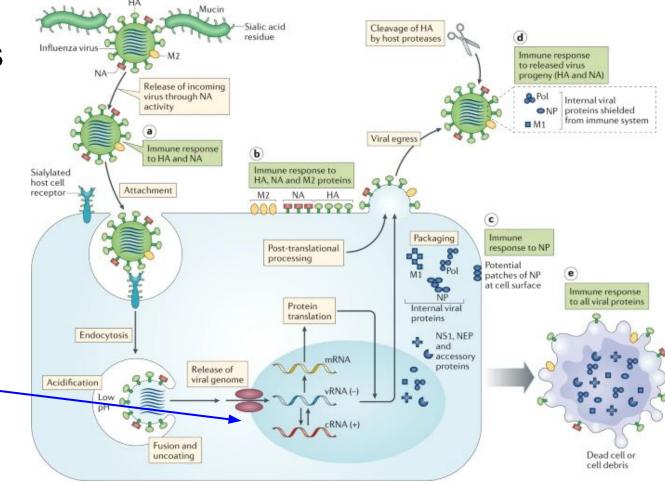


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Viral contents are released and viral RNA is reproduced, with the help of host components

Krammer. The human antibody response to influenza A virus infection and vaccination, 2020.

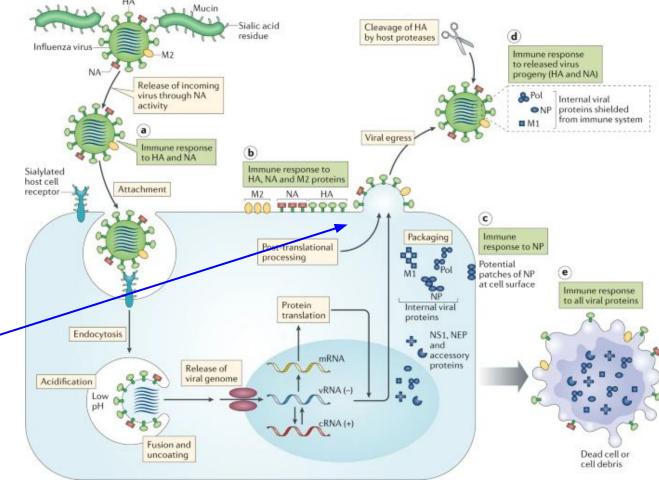


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New viruses are assembled and leave the cell through viral egress

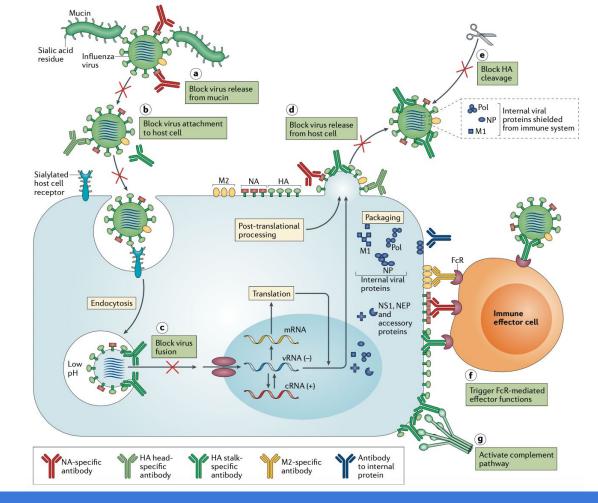
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Krammer. The human antibody response to influenza A virus infection and vaccination, 2020.

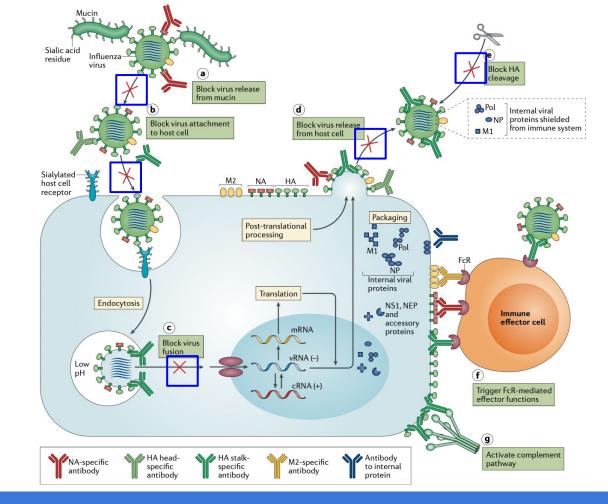


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Mechanisms for viral treatments include antibody binding to viral proteins that block / disrupt steps in the viral replication process

Krammer. The human antibody response to influenza A virus infection and vaccination, 2020.

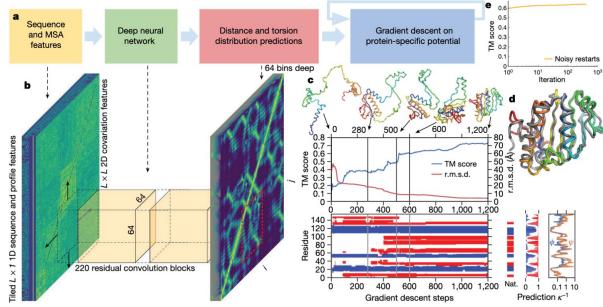


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AlphaFold

- Protein structure prediction: determining the 3D shape of a protein from its amino acid sequence
- Based on neural network that predicts distances between pairs of residues, then energy minimization to determine most likely 3D shape



Senior et al. Improved protein structure prediction using potentials from deep learning, 2020.

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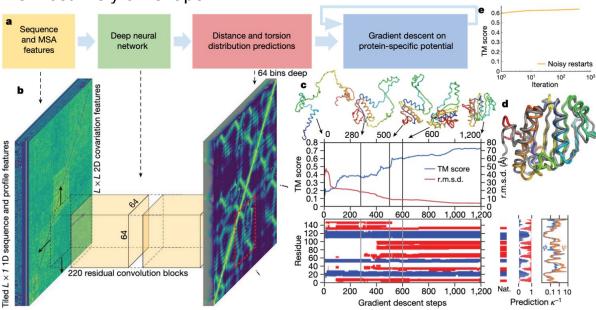
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AlphaFold

- Protein structure prediction: determining the 3D shape of a protein from its amino acid sequence
- Based on neural network that predicts distances between pairs of residues, then energy minimization to determine most likely 3D shape

Used approach to release structure predictions for proteins associated with SARS-CoV-2

Senior et al. Improved protein structure prediction using potentials from deep learning, 2020.



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Beck et al.

- Deep learning-based drug-target interaction model that predicts whether commercially available drugs can act on viral protein of SARS-CoV-2
- Extracted amino acid sequences of proteins from the SARS-CoV-2 replication complex
- Can use sequence models from NLP to model the data!

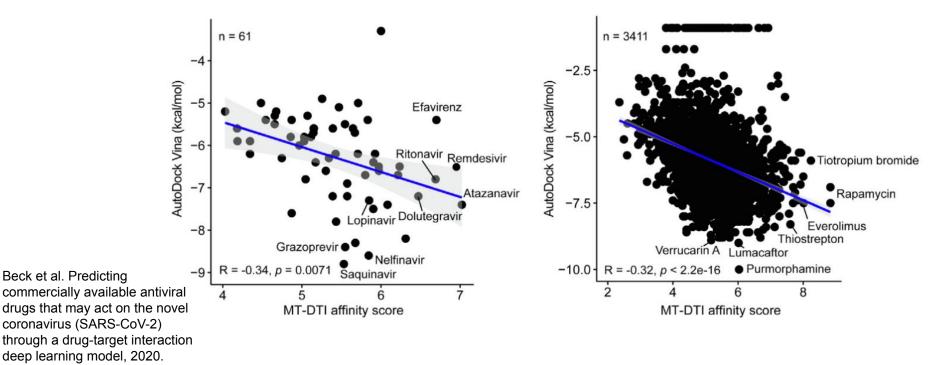
Atazanavir	COC(=O)NC(C(=O)NC(Cc1ccccc1)C(O)CN(Cc1ccc(-c2ccccn2)cc1)NC(=O)C(NC(=O)OC)C(C)(C)C)C(C)(C)C)C(C)C(C)C(C)C(C)C
Remdesivir*	CCC(CC)COC(=0)[C@H](C)N[P@](=0)(OC[C@@H]1[C@H]([C@H]([C@](O1)(C#N)C2 = CC = C3N2N = CN = C3N)O)O)
	OC4 = CC = CC = C4
Efavirenz*	O = C1Nc2ccc(Cl)cc2[C@@](C#CC2CC2)(C(F)(F)F)O1
Ritonavir	CC(C)c1nc(CN(C)C(=0)NC(Cc2cccc2)CC(0)C(Cc2cccc2)NC(=0)OCc2cncs2)C(C)C)cs1
Dolutegravir	CC1CCOC2Cn3cc(C(=0)NCc4ccc(F)cc4F)c(=0)c(0)c3C(=0)N12
Asunaprevir	C = CC1CC1(NC(=0)C1CC(Oc2ncc(OC)c3ccc(C1)cc23)CN1C(=0)C(NC(=0)OC(C)(C)C)C(C)(C)C)C(=0)NS(=0)(=0)C1CC1CCC(C)CCCCCCCCCCCCCCCCCCCCCCCCCCC
Ritonavir*	CC(C)c1nc(CN(C)C(=0)N[C@H](C(=0)N[C@@H](Cc2cccc2)C[C@H](0)[C@H](Cc2cccc2)NC(=0)OCc2cncs2)C(C)C)cs1
Simeprevir*	$COc1ccc2c(0[C@H]3CC4C(=0)N(C)CCCC/C = C \ [C@H]5C[C@@]5(C(=0)NS(=0)(=0)C5CC5)NC(=0)[C@@H]4C3)cc(-c3nc(C)CCCC) \ (C)CCCC/C = C \ (C)CCCCC/C = C \ (C)CCCC/C = C \ (C)CCCCC/C = C \ (C)CCCC/C = C \ (C)CCCCC/C = C \ (C)CCCC/C = C \ (C)CCCC/C = C \ (C)CCCCC/C = C \ (C)CCCCC/C = C \ (C)CCCCC/C = C \ (C)CCCCCC/C = C \ (C)CCCCCC/C = C \ (C)CCCCCC/C = C \ (C)CCCCCC/C = C \ (C)CCCCCCCC/C = C \ (C)CCCCCCC/C = C \ (C)CCCCCCC/C = C \ (C)CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC$
	(C)C)cs3)nc2c1C

Beck et al. Predicting commercially available antiviral drugs that may act on the novel coronavirus (SARS-CoV-2) through a drug-target interaction deep learning model, 2020.

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Beck et al.

Identified promising drugs such as atazanavir, remdesivir, and others

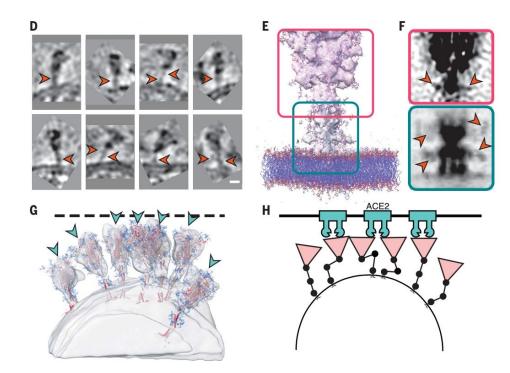


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CryoET high resolution imaging of virus spike structure

- CryoET (cryogenic electron tomography) imaging can provide high-resolution visualization of virus particles
- Analysis of virus spike (surface protein) structure, organization and variability can provide insight into how it binds to host cell receptors



Lecture 11 - 43

Turonova et al. In situ structural analysis of SARS-CoV-2 spike reveals flexibility mediated by three hinges, 2020.

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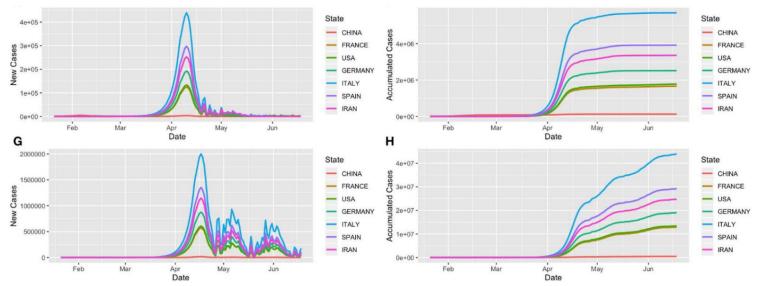
Additional application areas

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Epidemiology and disease forecasting

- Epidemiological models e.g., SIR, SEIR, based on numbers of susceptible, exposed, infected, recovered individuals help with anticipating and preparing for upcoming challenges
- Also efforts at deep learning-based forecasting of future cases



Hu et al. Forecasting and Evaluating Multiple Interventions for COVID-19 Worldwide, 2020. Shinde et al. Forecasting Models for Coronavirus Disease (COVID-19): A Survey of the State-of-the-Art, 2020.

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Telehealth

- Demand for virtual visits are up more than 1000x in some care settings¹
- Digital setting offers opportunity for AI algorithms
- Already many AI applications and even products popping up around use cases such as triaging, vitals measurement, and even cough analysis



¹https://www.aarp.org/health/con ditions-treatments/info-2020/teleh ealth-surges-during-coronavirusoutbreak.html Figure credit:

https://www.healthwise.org/blog/ patient-ed-telehealth-amid-covid-19.aspx

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Summary

Today we covered:

- Applications of AI in Healthcare through the lens of COVID-19
 - Al interpretation of chest radiology images
 - Modeling patient outcomes using EHR data
 - Finding treatments for the disease
 - Additional application areas

Next time:

- Unsupervised Learning and Reinforcement Learning

