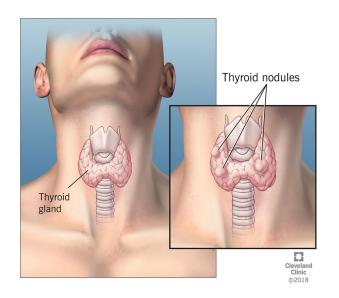


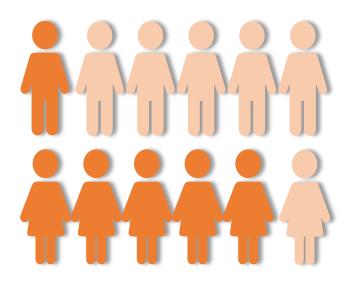


Background of thyroid nodules

In epidemiological surveys, about 4%-7% of the population have palpable thyroid nodules, compared with 17%-46% on ultrasound, and nearly 60% of the population have thyroid nodules reported in autopsy findings.

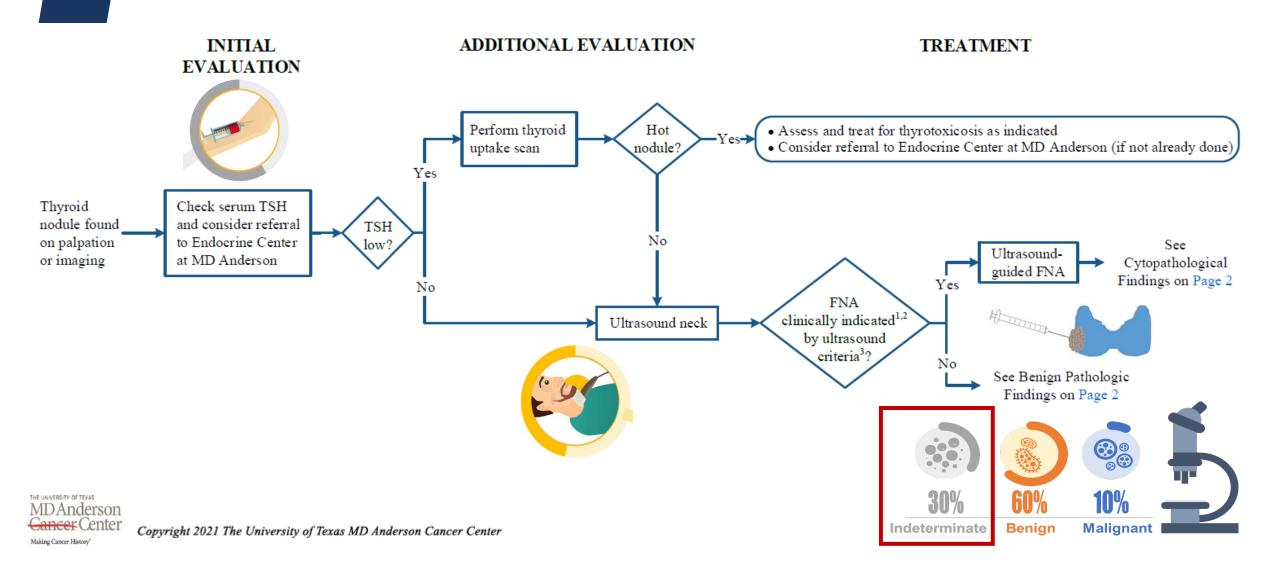
Despite of high incidence of thyroid nodules, only 7-15% of them are malignant.





- [1] Fagin JA & Wells SA, Jr. (2016). N Engl J Med 375, 1054-1067
- [2] Li Y, et al. (2021). Front Endocrinol (Lausanne) 12, 676144
- [3] Zhou J, et al.(2020) Endocrine 70, 256-279
- [4] K. D. Burman, L. Wartofsky. (2015). N Engl J Med 373, 2347-2356
- [5] C. Durante et al. (2018). JAMA 319, 914-924

Clinical evaluation of patients with thyroid nodules



Nucleic acid-based molecular testing for thyroid nodules

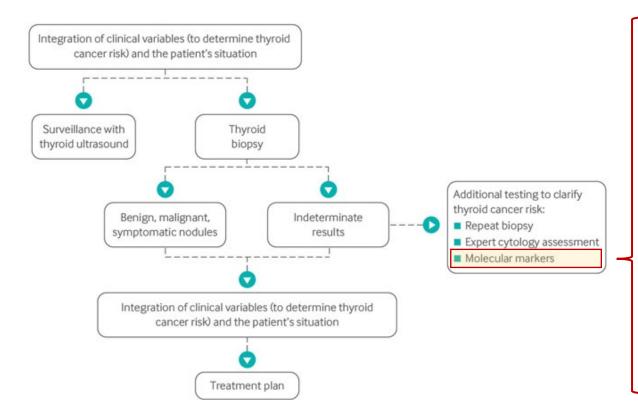


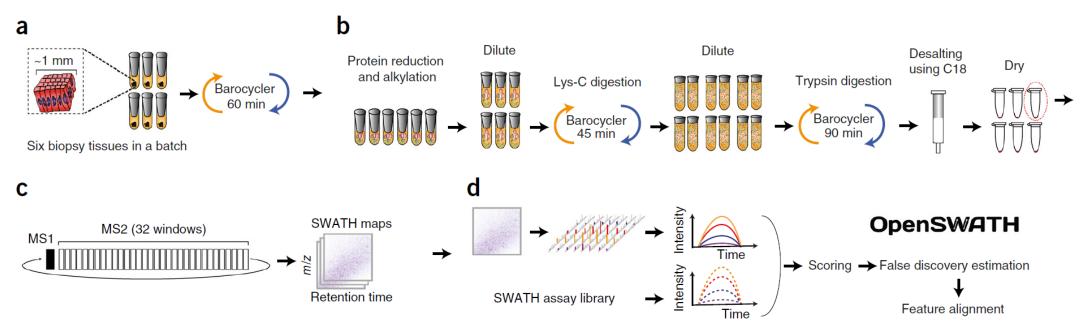
Table 1 | Commercially available molecular tests for cytologically indeterminate thyroid nodules

lable 1 Colliner		motecutar	tests	ror cy	tologi	catty in	ueterminate	tnyroid nodules				
Test	Cytology (Bethesda)	Number of cases	SN	SP	PPV	NPV	Test result	Clinical utility	Refs			
GEC	III	129	90	53	38	95	Benign	Active surveillance	19			
							Suspicious	Surgery ^a	19			
	IV	81	90	49	37	51	have demon to positive p	te thyroid nodules ¹ strated significant redictive values repo	variatio orted in			
ThyroSeqv2	III	96	91	92	77	97	pathology in GEC-suspicious nodules the sensitivity (83–100%) and specificity (GEC testing vary across studies, but ma					
	IV	143	90	93	83	96	lacked long-term follow-up of GEC-benig and/or histopathological correlation of C ings ^{17,19,22,23} . Overall, the low specificity ex use as a rule-in test, and thyroid lobectomy,					
ThyroSeq v3	III	154	91	85	64		mum, should be performed for diagnostic p GEC-suspicious thyroid nodules (TABLE 1).					
	IV	93	97	75	68	98	Negative	Active surveillance	25			
							Positive	Surgery ^a	25			
RosettaGX Reveal	III and IV	150	74	74	43	92	Benign	Active surveillance	113			
							Suspicious	Surgery ^a	113			
ThyGenX or ThyraMIR	III	58	94 80	80 68	68	97	Negative	Surgery or active surveillance	114			
							Positive	Surgery ^a	114			
	IV	51	82	91	82	91	Negative	Surgery or active surveillance	114			
							Positive	Surgery ^a	114			

GEC, gene expression classifier testing; NPV, negative predictive value; PPV, positive predictive value; SN, sensitivity; SP, specificity.
*Extent of surgery (thyroid lobectomy or total thyroidectomy should be determined by clinical variables, ultrasonographic features and patient preference).

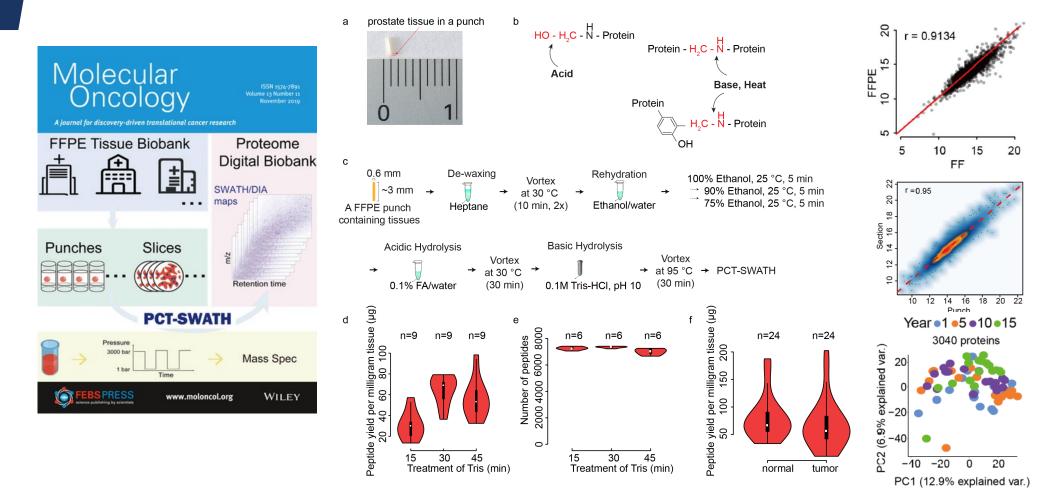
[1] Singh Ospina N, *et al.* (2020). BMJ 368, I6670 [2] Wang TS & Sosa JA (2018). Nat Rev Endocrinol 14, 670-683

Pressure cycling technology assisted sample preparation



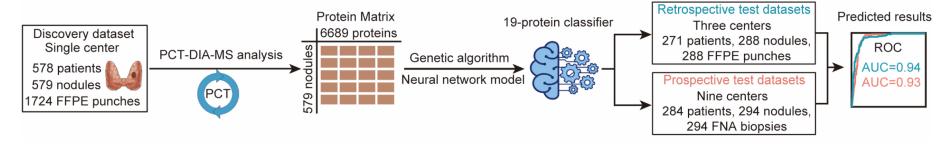
- 1. High-throughput
- 2. Small amount of samples(<1mg tissue weight)
- 3. Fast (100s-1000s)
- 4. Robustness
- 5. Potential for quantification of entire proteome
- [1] Gillet LC, ..., Bonner R & Aebersold R (2012). Mol Cell Proteomics 11, O111.016717
- [2] Guo T, ..., Jochum W & Aebersold R (2015). Nat Med 21, 407-413
- [3] Powell K (2018) Technology to watch in 2018. Nature 553, 531-534

PCT assisted sample preparation from FFPE tissues



Zhu Y, ..., Aebersold R & Guo T (2019). Mol Oncol 13, 2305-2328

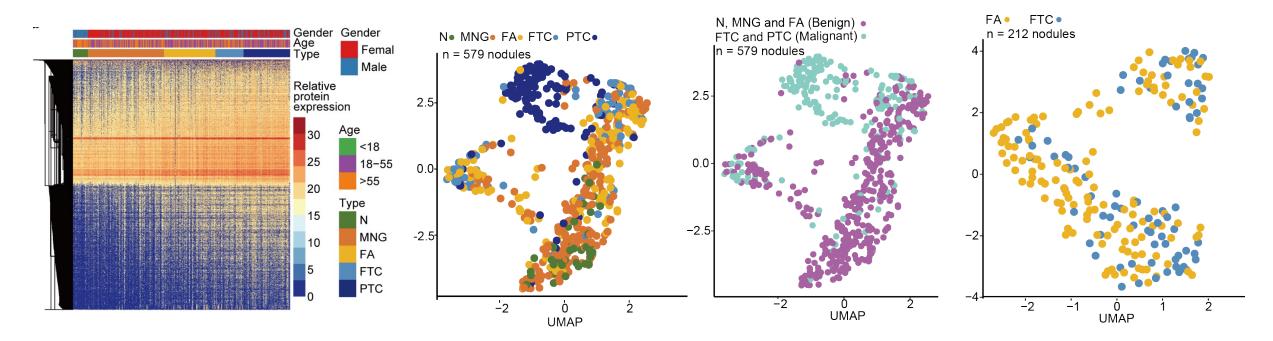
Study design and patient characteristics



Clinico-pathologic characteristics of the study cohorts

			Independent		
		Discovery dataset (FFPE)	Retrospective test datasets (FFPE)	Prospective test datasets (FNA)	All
Total no.					
	Clinical centers	1	3	9	12°
	Patients	578	271	284	1133
	Nodules	579	288	294	1161
	FFPE cores	1724	288	0	2012
	Fine needle asperation biopsies	0	0	294	294
	DIA files	1780	576	294	2650
Histopathology diagnosis					
	Normal tissues / lymphocytic thyroiditis	s 40 (6.9%)	16 (5.6%)	6 (2.7%)	64 (5.5%)
	Moltinodular goiter	203 (35.1%)	44 (15.3%)	62 (21.1%)	309 (26.6%)
	Follicular adenoma ^a	137 (23.7%)	84 (29.2%)	23 (7.8%)	244 (21.0%)
	Follicular thyroid carcinoma ^a	75 (13.0%)	52 (18.1%)	4 (1.4%)	131 (11.3%)
	Papillary thyroid carcinoma	124 (21.4%)	92 (31.9%)	197 (67.0%)	413 (35.6%)
Bethesda classification	า				
	1	-	-	32 (10.9%)	-
	II	-	-	41 (13.9%)	-
	III	-	-	52 (17.7%)	-
	IV	-	-	22 (7.5%)	-
	V	-	-	20 (6.8%)	-
	VI	-	-	127 (43.2%)	-

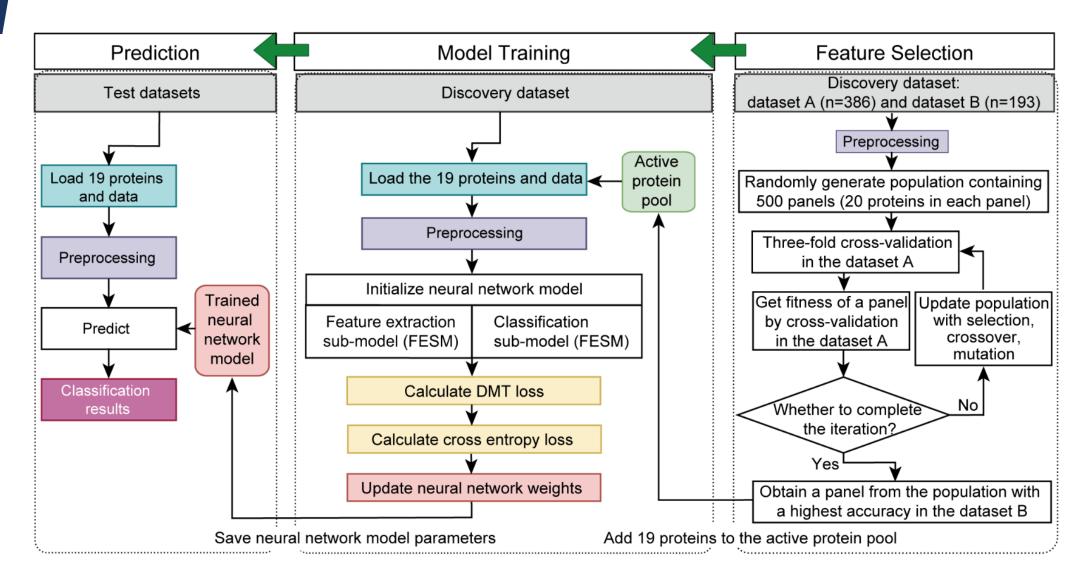
Global thyroid proteome profile



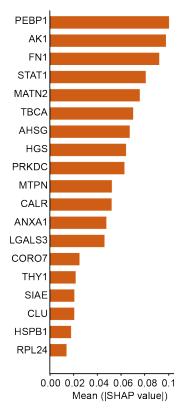
579 thyroid tissue specimens
5312 proteins (rows) are clustered
without supervision
Samples (columns) are ordered
based on the tissue types

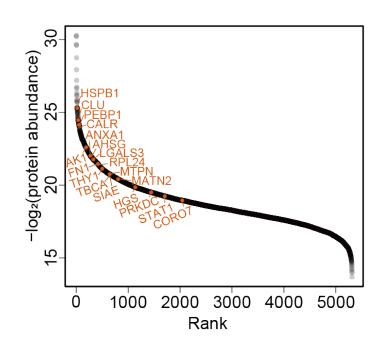
UMAP plots showing global snapshots comparing the indicated types of thyroid tissues using 5312 proteins for all subtypes; malignant versus benign nodules;FA versus FTC.

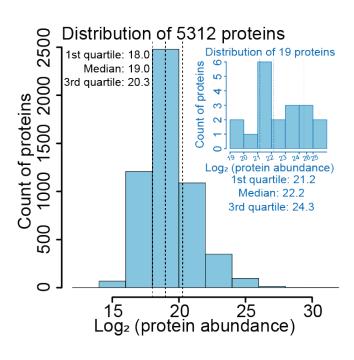
Schematic of principal classifier model



19 selected protein features



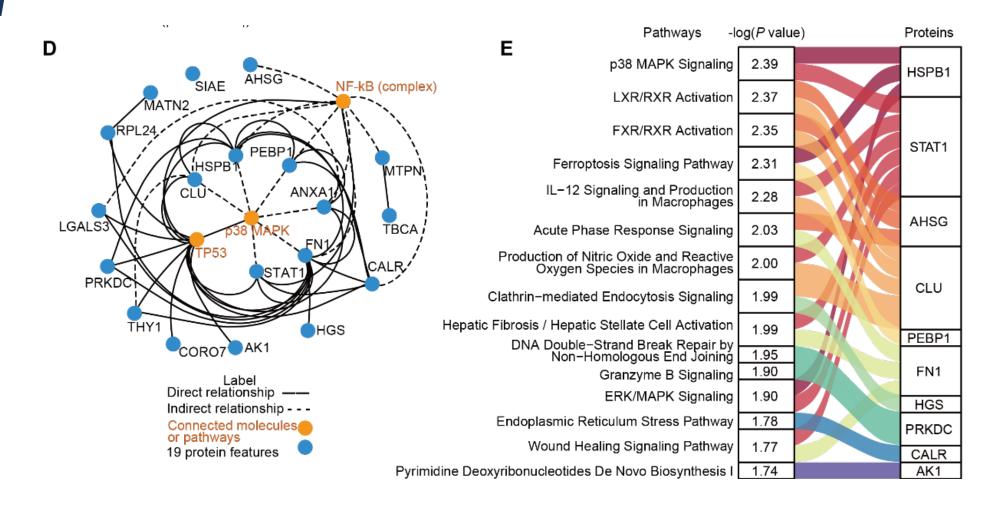




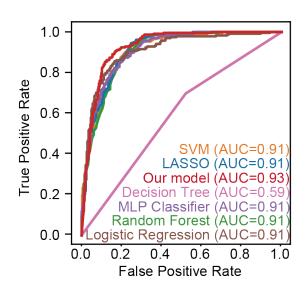
The importance rank of the selected 19 protein features was interpreted by SHapley Additive exPlanations (SHAP) algorithm.

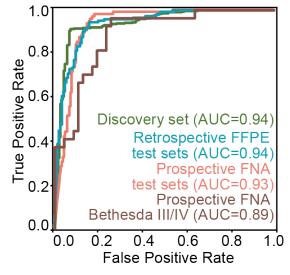
Protein abundance distribution of the 19 features

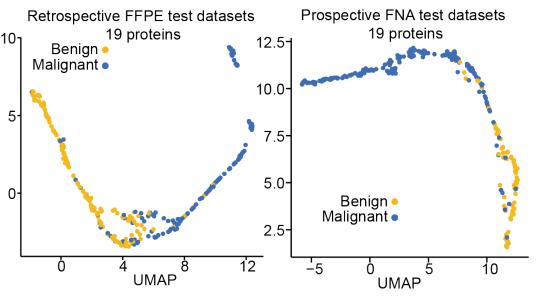
Biological insights on 19 selected features



Multicenter Clinical Evaluation of patients with thyroid nodules







Compared with other six different machine models of 19 selected features

Retrospective sets FFPE, 3 hospitals, 288 nodules

Prospective sets FNA, 9 hospitals, 294 nodules

Multicenter Clinical Evaluation of patients with thyroid nodules

Performance of ThyroProt classifier

Method	Prevalence	Sensitivity	Specificity	NPV	PPV	Accuracy
ThyroProt_FFPE	50.00%	84.03%	93.75%	93.08%	85.44%	88.89%
ThyroProt_FNA	68.03%	91.50%	71.28%	87.14%	79.76%	85.03%
ThyroSeq V3 ¹	29.57%	93.42%	81.22%	96.71%	67.62%	84.82%
GSC ²	23.68%	91.11%	68.28%	96.12%	47.13%	73.68%
GEC ³	32.08%	91.76%	51.67%	93.00%	47.27%	64.53%

	N/L	MNG	FA	FTC	PTC	All
Discovery set Retrospective FFPE test sets Prospective FNA test sets All	40/40	188/203	116/137	62/75	122/124	528/579
	15/16	43/44	77/84	37/52	84/92	256/288
	1/8	49/63	17/23	2/4	181/196	250/294
	56/64	280/310	210/244	101/131	387/412	1034/1161
	0 5			75	5%	100%

^[1] Steward, David L., et al. *JAMA oncology*, 2019

^[2] Kepal N. Patel., et al. JAMA Surgery, 2018

^[3] Alexander, Erik K., et al. *New England Journal of Medicine*, 2012

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