

# Knowledge Diffusion Through Networks

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# How is Knowledge Diffusion Shaped by Social Networks?

- Frictions that impede knowledge flows have the strong potential to affect efficiency and inequality— for example,
  - Slow diffusion of cost-reducing innovation (e.g. Griliches 1957)
  - Deviations from the law of one price (e.g. Jensen 2007)
  - Possible herding on inefficient choice (Banerjee 1992)→ Motivates policies that facilitate flows of knowledge
- Understanding the *structure* of knowledge frictions is key to implementing such ‘information’ policies well
- This paper: study a medical context. Doctors are positioned within a network and make prescription choices based on current knowledge
  - Provide evidence of doctor scripting behavior converging over time
  - Develop a social learning model that rationalizes this fact (no causal identification strategy)
  - Compare estimated vs. model-generated knowledge paths, inspect the mechanism, and perform counterfactuals

# Main Findings (so far)

- 1 Two Descriptive Statistics:
  - Fact #1: Prescription shares are converging across doctors
  - Fact #2: Convergence rates are increasing in network centrality
  - These facts are also evident in generic substitution context
    - Not just about advertising
  - Both facts are robust to different measures of centrality and convergence
- 2 Imposing some additional structure, we develop an econometric approach to estimate doctor-specific knowledge paths
- 3 We develop a dynamic model of learning on a network
  - We estimate initial knowledge stocks and network structure
  - Model-generated learning matches estimated paths

# Related Literature

## 1. Theory papers on dynamics of social learning

- Banerjee (1992), Smith and Sorensen (2008), Acemoglu et al (2011), etc.
- Our contribution:
  - ① A quantitative paper
  - ② Transition dynamics
  - ③ Social learning on a fairly general estimated network

## 2. Idea diffusion papers

- Grossman-Helpman (1991), Luttmer (2007), Lucas (2009), Lucas-Moll (2013), Perla-Tonetti (2013), Buera-Oberfield (2017), etc.
- Our contribution:
  - ① Empirical focus with micro data and specific tangible technologies
  - ② Incorporate geography and other elements of networks (not random search)
  - ③ Bayesian learning instead of  $\max(z, z')$

## Related Literature

### 3. Learning in pharmaceutical markets

- Erdem and Keane (1996), Akerberg (2003), Crawford and Shum (2005), Arrow, Bilir and Sorensen (2017), Dickstein (2018)
  - Approach: dynamic discrete choice estimation with Bayesian learning
  - Estimate importance of 'signals' from network neighbors
- Our contribution: prescription convergence depends on network (endogenous signal structure)

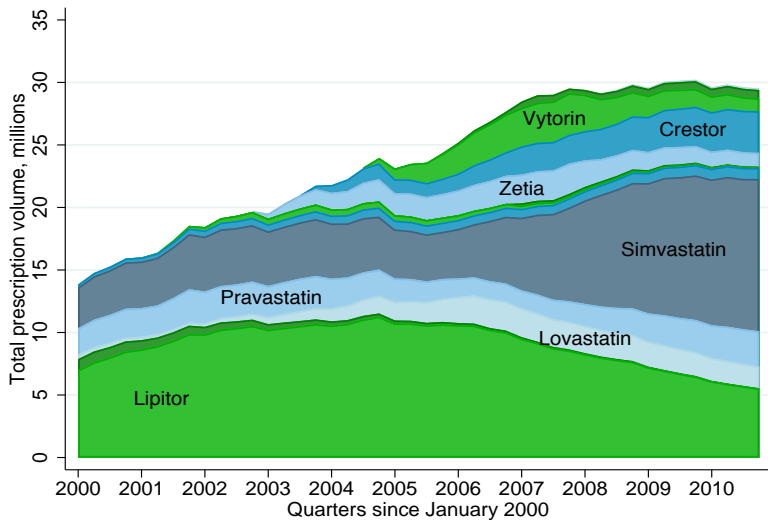
### 4. Papers measuring healthcare disparities: Dartmouth Atlas (Wennberg et al 1996, Munson et al 2013) and Cooper et al (2015)

- Wide variation in healthcare quality, efficiency across regions (e.g. high vs. low generic prescription share)
- Explained by unobserved differences in patient types or in knowledge?
- Our contribution: our network-learning mechanism can explain observed convergence—*changes* in the extent of medical care disparities
  - Central doctors learn faster; peripheral doctors slower
  - This mechanism can rationalize treatment disparities

## Data: Prescriptions from IMS Health (IQVIA, Xponent)

- We observe prescriptions by doctor, drug, and month
- Our data cover January 2000 through December 2010
  - This time period covers sequence of 12 drug innovations
  - Key entrants: three generics, three new 'molecular entities'
  - Today, within-spell analysis. In future, introduction of new drugs.
- Includes all U.S. doctors with at least 10 cholesterol drug prescriptions in *both* 2000 and 2010
  - 131,323 doctors  $\times$  132 months  $\times$  up to 18 drugs
  - We observe location (five-digit zipcode) for all doctors
  - For  $\sim$  47K, medical school, cohort, specialty also observed
- Extraordinarily precise data on repeated technology adoption and prescription decisions by the universe of relevant individuals
- Unfortunately, no data on patients or insurance plans

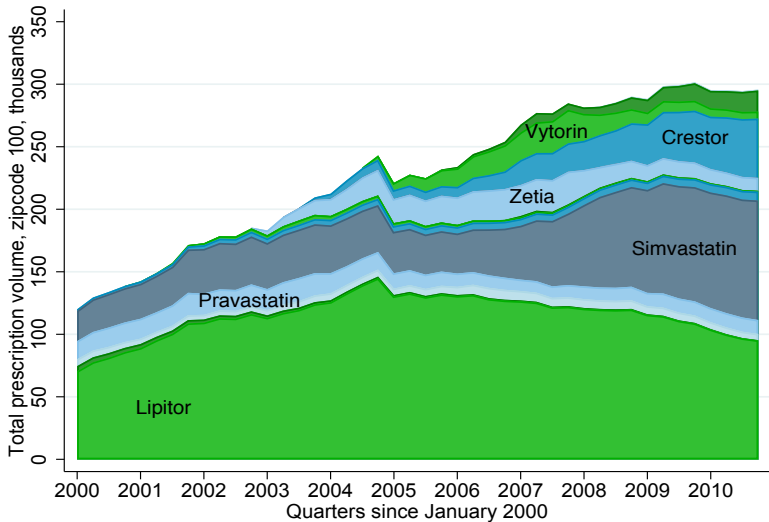
# Aggregate Evolution in Prescribing, Jan. 2000–Dec. 2010



→ Suggests changes in perceived drug 'qualities'

# Evolution in Prescribing: NYC

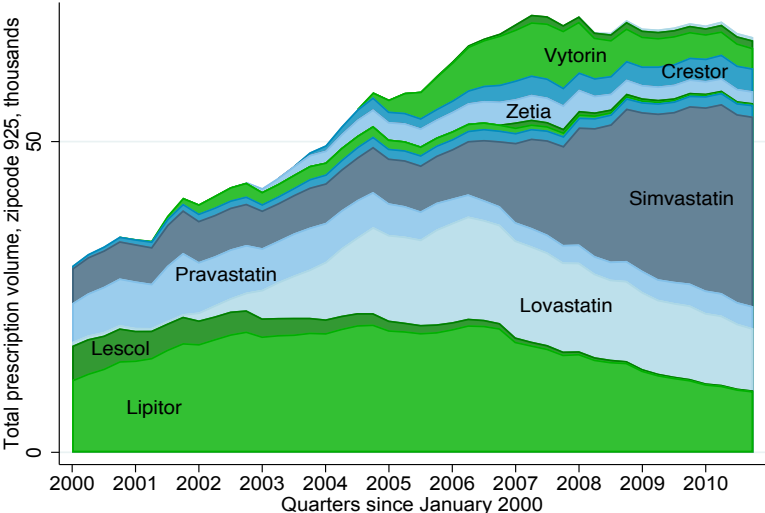
→ Variation and evolution is different for areas like New York, NY...





# Evolution in Prescribing: Hemet, CA

→ ...Relative to remote areas like Hemet, CA



## Within-molecule Substitution to Generic also varies

- Gradual diffusion is observed even for generics
- Generic substitution is only partial six months after generic release, but is essentially complete by December 2010 (final month)

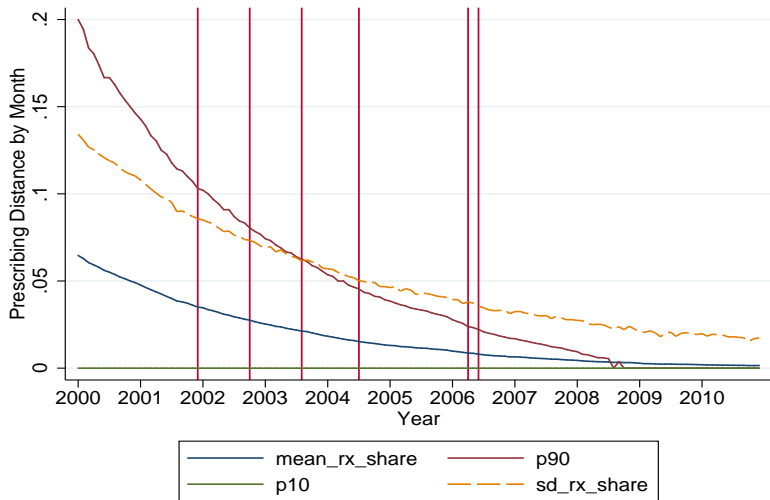
	<u>Generic Share in Prescriptions</u>		
	Lovastatin	Pravastatin	Simvastatin
<u>After six months</u>			
Mean	0.8280	0.8197	0.8616
St Dev	0.3382	0.2793	0.2079
5 <sup>th</sup> Percentile	0	0	0.448
95 <sup>th</sup> Percentile	1	1	1
 <u>December 2010</u>			
Mean	0.9995	0.9930	0.9970
St Dev	0.0188	0.0588	0.0276
5 <sup>th</sup> Percentile	1	1	0.995
95 <sup>th</sup> Percentile	1	1	1

## Different Types of Convergence

- Will summarize doctor scripting patterns by drug shares
- Could have all doctors moving steadily towards same final-period drug shares
  - Should see decreasing distance between average share and final share over time
- Could have doctors drug-shares becoming more similar to each other each period, but big fluctuations in those drug shares across periods
  - Should see decreasing variation in drug shares across doctors over time.
- Reminiscent of  $\beta$  and  $\sigma$  convergence in growth regression literature
- Will now show pictures that show within-spell both types of convergence are present

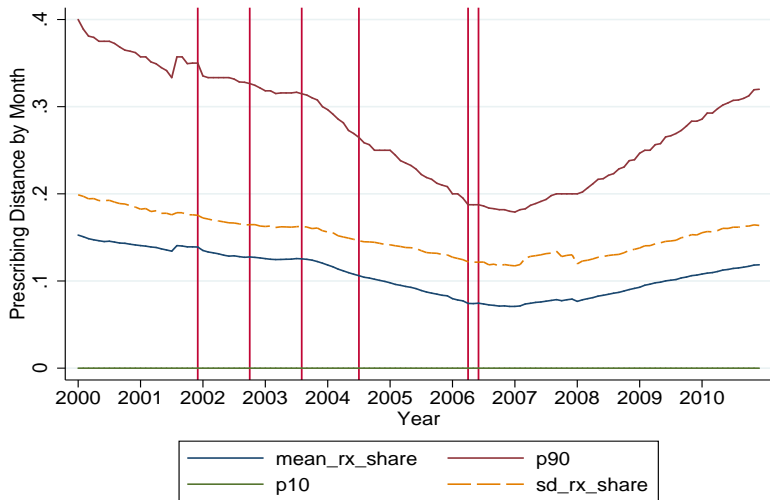
# Fact #1: Prescriptions are converging across doctors

Lescol



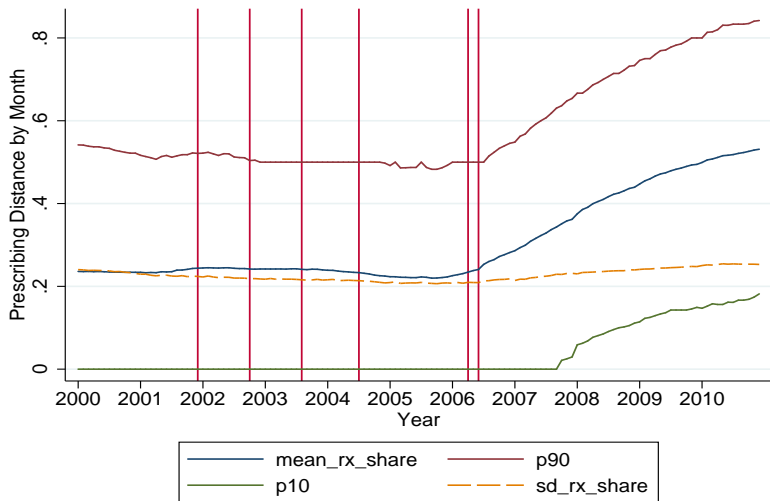
# Fact #1: Prescriptions are converging across doctors

## Pravastatin



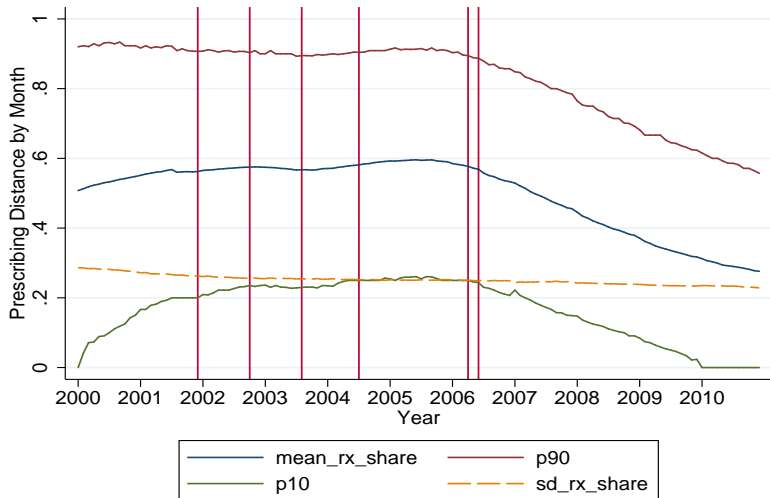
# Fact #1: Prescriptions are converging across doctors

Simvastatin



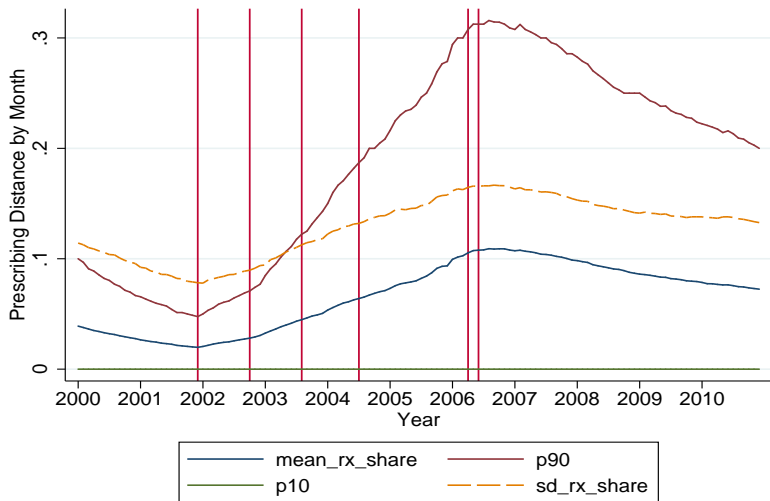
# Fact #1: Prescriptions are converging across doctors

Lipitor



# Fact #1: Prescriptions are converging across doctors

Lovastatin





## Fact #1: Prescriptions are converging across doctors

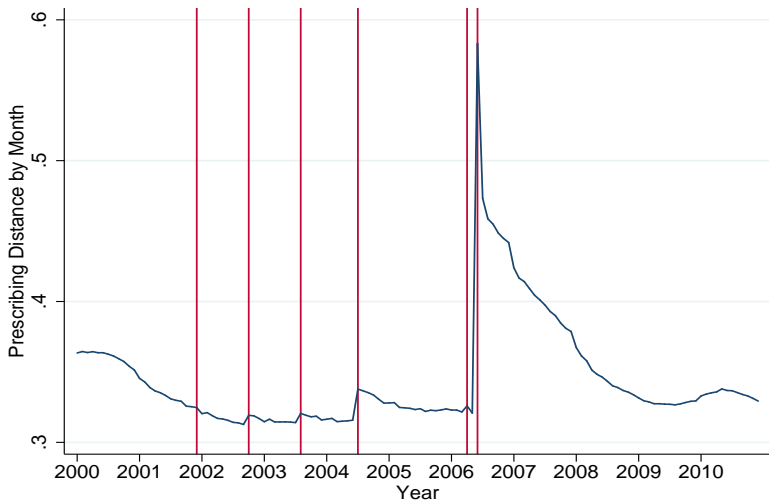
- Consider drugs  $\mathcal{D}_t = \{1, 2, \dots, D_t\}$  available in month  $t$
- Group months into spells defined by major drug introductions
- $\pi_{idt}$  is drug- $d$  share of doctor  $i$ 's month- $t$  prescriptions

Distance measure  $y_{it}$ : Euclidean prescription distance between  $i$  at  $t$  and average doctor at end of spell  $T(t)$ ,

$$y_{it} \equiv \left( \sum_{d=1}^{D_t} \left( \pi_{idt} - \bar{\pi}_{dT(t)} \right)^2 \right)^{\frac{1}{2}}$$

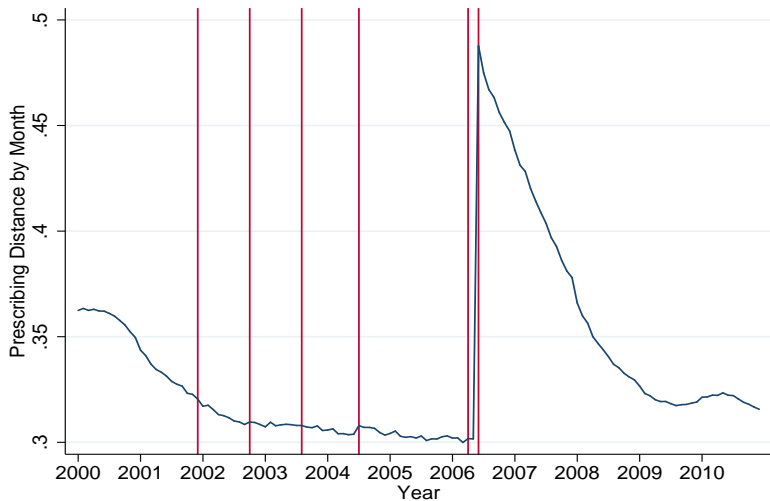
- $\bar{\pi}_{dT(t)}$  is average prescription vector at end of current spell
- Changes in  $\mathcal{D}_t$  can affect convergence mechanically; perform analysis using a) all drugs, b) six molecules available at  $t = 0$
- Start with simple plots of  $\bar{y}_t = \frac{1}{I} \sum_i y_{it}$  over time

## Scripting Patterns Converge across Doctors



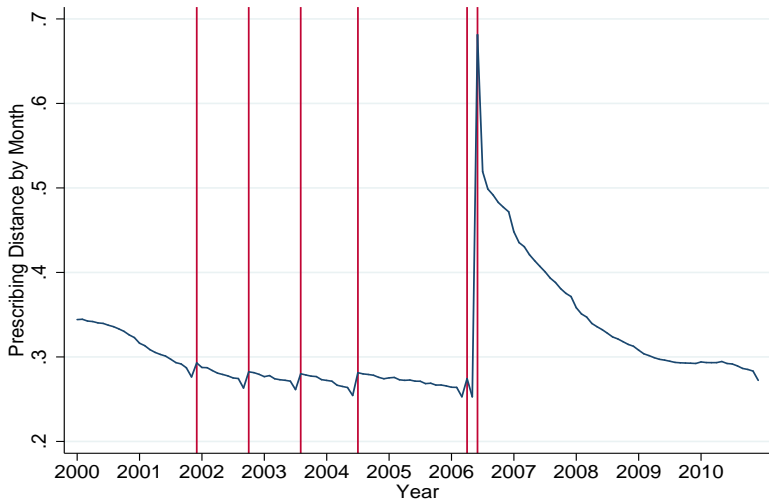
# Scripting Patterns Converge across Doctors

Same convergence pattern for initial 6 drugs



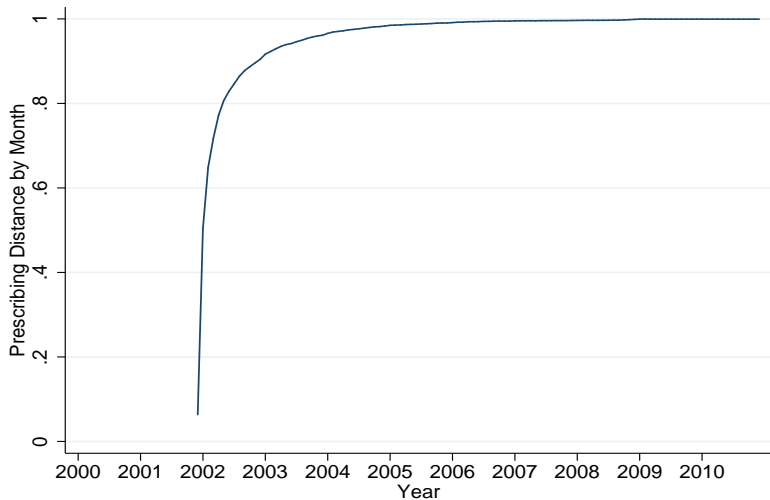
# Scripting Patterns Converge across Doctors within Zipcode

Same convergence pattern for initial 6 drugs



# Generic Prescription Share Converges

→ Within-molecule generic substitution, Lovastatin



## Fact #2: Central doctors converge faster

- 1 Measure simple doctor- $i$  specific convergence rate  $\beta_i$

$$y_{it} = \beta_i \times t + \delta_i + \delta_t + \epsilon_{it}$$

- 2 Regress  $\beta_i$  on measure of the centrality of each doctor  
2.1 'Gravity' regression to construct centrality measure:

$$\tilde{X}_{ij} = \alpha_1 \widetilde{dist}_{ij} + \alpha_2 \widetilde{school}_{ij} + \alpha_3 \widetilde{cohort}_{ij} + \alpha_4 \widetilde{specialty}_{ij} + \delta_i + \delta_j + \epsilon_{ij}$$

where  $X_{ij} \equiv (\sum_d (\bar{\pi}_{id} - \bar{\pi}_{jd})^2)^{1/2}$ ,  $\tilde{Z}_{ij} = 1/(1 + Z_{ij})$

- Estimate using a random sample of  $\sim 3,200$  doctors
- Medical specialty is dominated by three groups: internal medicine (36%), family practice (28%), and cardiology (10%)
- $\sim 250$  medical schools, 8.5K zipcodes
- Median doctor graduates in 1981

## Fact #2: Central doctors converge faster

2.2 Adjacency matrix  $A \equiv [A_{ij}]$ :

$$A_{ij} \equiv \exp\{\hat{\alpha}_1 \widetilde{dist}_{ij} + \hat{\alpha}_2 school_{ij} + \hat{\alpha}_3 cohort_{ij} + \hat{\alpha}_4 specialty_{ij}\}$$

2.3 Centrality: the eigenvector  $c \equiv [c_i]$  associated with the largest eigenvalue of  $A$  contains the centrality index for each doctor  $i$

2.4 Regress doctor-specific convergence rate  $\beta_i$  on centrality index  $c_i$

$$\hat{\beta}_i = \lambda c_i + x_i + \eta_{z(i)} + \epsilon_i,$$

where  $x_i$  is prescription volume, advertising,  $\eta_{z(i)}$  are zipcode FE

# 'Gravity' Estimates— Bilateral prescription 'proximity'

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Proximity ( $\widetilde{dist}$ )	.1228 <sup>a</sup> (.0013)	.1175 <sup>a</sup> (.0017)
Same medical school	.0036 <sup>a</sup> (.0002)	.0038 <sup>a</sup> (.0002)
Same medical school cohort	.0004 <sup>a</sup> (.00003)	.0003 <sup>a</sup> (.00003)
Same medical specialty	.0026 <sup>a</sup> (.00004)	.0025 <sup>a</sup> (.00004)
Same school $\times \widetilde{dist}$		-.0281 <sup>a</sup> (0073)
Same cohort $\times \widetilde{dist}$		.0184 <sup>a</sup> (.0028)
Same specialty $\times \widetilde{dist}$		.0054 <sup>c</sup> (.0029)
Doctor $i$ FE	Y	Y
Doctor $j$ FE	Y	Y
$R^2$	0.6338	0.6338
Observations	10220788	10220788

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## Convergence rate is increasing in centrality index

- Equation:  $\hat{\beta}_i = \lambda c_i + x_i + \eta_{z(i)} + \epsilon_i$

	Rx Distance		Generic Share	
Centrality	-.3957 <sup>a</sup> (.0436)	-.3748 <sup>a</sup> (.0436)	.0324 <sup>a</sup> (.0090)	.0322 <sup>a</sup> (.0090)
Rx volume	-.0519 <sup>a</sup> (.0190)	-.0073 (.0154)	-.0141 <sup>a</sup> (.0032)	-.0145 <sup>a</sup> (.0033)
Advertising		-.1290 <sup>a</sup> (.0110)		.0012 (.0022)
Zipcode FE	Y	Y	Y	Y
R <sup>2</sup>	.2462	.2497	.2039	.2039
Observations	37417	37417	35907	35907

Robustness: a) degree centrality, b) estimate gravity coefficients using excluded physician sample or observations from excluded time period

## Summary of descriptive results

- Fact #1: Prescription shares are converging across doctors
- Fact #2: Convergence rates are increasing in network centrality
- These facts are also evident in generic substitution context
- Both facts are robust to alternative measures of convergence and centrality

Next steps motivated by descriptive results:

- ① Develop structural model of scripting and learning
- ② Use structural equations to estimate doctor-specific learning rates, network structure, and initial knowledge
- ③ Counterfactuals in simulated environment

# Why a Structural Model?

- With structural model, can see how lowering barriers to knowledge diffusion will affect learning, accounting for endogeneity of effort
- Quantify how much barriers to knowledge affect drug mis-prescribing
  - Could connect to the value of improving doctors' knowledge (continuing ed., electronic medical records with or without decision support tools, etc.)
- Quantify which variables most affect knowledge diffusion: geography, age, common med school, etc. (analogy to distance, common border, common language, etc. in trade)
  - Which doctors would it be most useful to educate? Those who know least? Those most connected? Probably both important, but not one in the same, so meaningful tradeoff.
- Use detailed data as model selection guide: different models of networks and learning

## Roadmap: Summary of Structural Model

- Drug discrete choice problem relates drug-shares  $\pi_{idt}$  to beliefs about drug quality and risk aversion (static problem given beliefs)
- Beliefs are distributed Normal and update according to Bayesian learning
  - LoM for beliefs depends on number of signals and signal distribution
- Social learning: Signals received depend on network
  - Number of signals depends on who doctor is connected to and how much those doctors know
  - Signal distribution depends on who doctor is connected to, how much those doctors know, and *what those doctors believe* (not today)
- Doctors choose investment in learning (number of signals) to maximize PDV of their expected utility (not today)
- No doctor-specific patient population and no advertising (as of now)

# Patient Utility and Doctor Beliefs

- Consider a spell with a fixed set  $\mathcal{D}$  of available drugs
- Each doctor  $i \in \{1, \dots, N\}$  treats a unit measure of patients  $v_i \in [0, 1]$  in period  $t = 1, 2, \dots, T$
- Patient  $v_i$  reward from drug  $d \in \mathcal{D}$  is

$$u_{dt}(\varepsilon_{dt}(v_i)) = \beta_d + \varepsilon_{dt}(v_i)$$

- The true unconditional efficacy of drug  $d$ ,  $\beta_d$
- An idiosyncratic observed shock,  $\varepsilon_{dt}(v) \sim \text{Gumbel } F(\cdot)$
- However, doctors are only imperfectly informed about  $\{\beta_d\}_{d \in \mathcal{D}}$ 
  - Doctor  $i$  beliefs at  $t$  about  $\beta_d$  are summarized by a Normal distribution  $G_{idt}(\cdot)$  with mean  $\tilde{\beta}_{idt}$  and variance  $\sigma_{idt}^2$

## Doctor Expected Utility from Treating Patient $v_i$

- Let doctors have CARA preferences represented by  $\mathcal{U}_{idt}(\varepsilon_{dt}(v_i), x)$
- Expected utility of doctor  $i$  treating patient  $v_i$  with drug  $d$  given beliefs  $x \sim G_{idt}(\cdot)$  is

$$\begin{aligned}\mathcal{U}_{idt}(\varepsilon_{dt}(v_i), \tilde{\beta}_{idt}, \sigma_{idt}) &:= \int -\exp(-\alpha(x + \varepsilon_{dt}(v_i))) dG_{idt} \\ &= -\exp(-\alpha\varepsilon_{dt}(v_i)) \int \exp(-\alpha x) dG_{idt} \\ &= -\exp(-\alpha\varepsilon_{dt}(v_i)) \exp\left(-\alpha\tilde{\beta}_{idt} + \frac{1}{2}\alpha^2\sigma_{idt}^2\right) \\ &= -\exp\left(-\alpha\tilde{\beta}_{idt} + \frac{1}{2}\alpha^2\sigma_{idt}^2 - \alpha\varepsilon_{dt}(v_i)\right)\end{aligned}$$

## Expected Utility, Beliefs, and Prescription Patterns

Assume learning is independent of prescription choice (verified later).

- Doctor  $i$  chooses a drug at  $t$  for each patient  $v_i$  to maximize expected doctor utility given current beliefs, yielding expected utility

$$U_{it}(\varepsilon_{dt}(v_i), \tilde{\beta}_{it}, \sigma_{it}) := \max_{d \in D} \{U_{idt}(\varepsilon_{dt}(v_i), \tilde{\beta}_{idt}, \sigma_{idt})\}.$$

- Given  $\varepsilon_{dt}(v_i)$  is distributed Gumbel, the doctor faces a standard multinomial choice problem, such that doctor  $i$  chooses drug  $d$  for patient  $v_i$  with probability  $\pi_{idt}$ :

$$\pi_{idt}(\varepsilon_{dt}(v_i), \tilde{\beta}_{it}, \sigma_{it}) :=$$

$$\Pr \{ \tilde{\beta}_{idt} - \alpha \sigma_{idt}^2 / 2 + \varepsilon_{idt}(v_i) > \tilde{\beta}_{id't} - \alpha \sigma_{id't}^2 / 2 + \varepsilon_{id't}(v_i), \forall d' \neq d \}$$
$$= \frac{\exp(\tilde{\beta}_{idt} - \alpha \sigma_{idt}^2)}{\sum_{d' \in D} \exp(\tilde{\beta}_{id't} - \alpha \sigma_{id't}^2)}$$

- Empirical analog:  $\pi_{idt}$  is drug- $d$  share in doctor  $i$ 's portfolio at time  $t$

## Doctor Expected Utility from Treating all Patients

- Finally, doctor- $i$ 's period payoff considering all the patients she treats at  $t$  is

$$W_{it}(\tilde{\beta}_{it}, \sigma_{it}) \equiv \int U_{it}(\varepsilon_{dt}(v_i), \tilde{\beta}_{it}, \sigma_{it}) dF(\varepsilon_{dt}(v_i)).$$

- And the dynamic problem of a doctor is to choose  $l_{it} \in \mathcal{R}^+$ :

$$V_{it}(\tilde{\beta}_{it}, \sigma_{it}) \equiv \max_{\{l_{i\tau}\}_{\tau=t}^{\infty}} \sum_{\tau \geq t} \delta^{\tau-t} (W_{i\tau}(\tilde{\beta}_{i\tau}, \sigma_{i\tau}) - c(l_{i\tau}))$$

L.O.M. for  $\beta_{it}$  and  $\sigma_{it}$  (which depends on  $l_{it}$ )



## Normal Bayesian learning: LoM for $\beta_{it}$ and $\sigma_{it}$ given $f_{it}$

- Prior beliefs are distributed around true values:  $\tilde{\beta}_{id0} \sim N(\beta_d, \sigma_d^2)$
- Each doctor receives  $f_{idt}$  signals at  $t$ , values  $\{x_{dn}\}$ ,  $n = 1, \dots, f_{idt}$ 
  - Signals accumulate into knowledge  $S_{idt} = S_{id0} + \sum_{u=1}^t f_{idu}$
  - For today, signal values are centered around truth:  $x_{dn} \sim N(\beta_d, \sigma_d^2)$
  - Let  $\bar{x}_{idt} = \sum_{n=1}^{f_{idt}} x_{dn} / f_{idt}$   
 $\implies \bar{x}_{idt} \sim \beta_d + \epsilon_{idt}$ , where  $\epsilon_{idt} \sim N(0, \sigma_d^2 / f_{idt})$
- Standard Bayesian learning then implies,

$$\tilde{\beta}_{idt+1} = \frac{\tilde{\beta}_{idt} S_{idt} + \bar{x}_{idt+1} f_{idt+1}}{S_{idt} + f_{idt+1}}$$

$$\sigma_{idt}^2 = \frac{\sigma_{id0}^2}{S_{idt}}$$

- Assumptions for today:
  - Assume signals are general information:  $f_{idt} = f_{it} \forall d, t$
  - Assume common prior knowledge:  $S_{id0} = S_{i0} \forall d$
  - Then  $S_{idt} = S_{it}$  and  $\sigma_{idt}^2 = \frac{\sigma_d^2}{S_{it}} \forall d, t$

## Knowledge diffusion through networks: $f_{it}$ and LoM for $S_{it}$

- Our model of  $i$ 's knowledge flow at  $t$  ( $f_{it}$ ) depends on (1) investment in learning  $l_{it}$ , (2) network connections, and (3) other doctors' stocks of knowledge as follows

$$f_{it+1} = l_{it} \times \sum_j \tau_{ij} S_{jt}$$

where  $\tau_{ij} \geq 0$  reflects the strength of network connections

- Then the law of motion for the stock of knowledge is

$$\begin{aligned} S_{it+1} &= Q(l_{it}, \mathcal{S}_t, \mathcal{T}) \\ &= S_{it} + f_{it+1} \\ &= S_{it} + l_{it} \times \sum_j \tau_{ij} S_{jt} \end{aligned}$$

- where  $\mathcal{S}_t$  is the vector of  $S_{it}$  and  $\mathcal{T}$  is matrix of  $\tau_{ij}$
- We've also worked with  $Q(S_{i,t}, \{l_{ijt}\}, \mathcal{T}) = \delta S_{i,t} + l_{it} \sum_j \tau_{ij}^{-\epsilon} l_{jt}$

## Summary of Model

- Doctor discrete choice relates drug-shares  $\pi_{idt}$  to beliefs  $\tilde{\beta}_{it}$ ,  $\sigma_{it}$  and risk aversion

$$\pi_{idt} = \frac{\exp(\tilde{\beta}_{idt} - \alpha\sigma_{idt}^2)}{\sum_{d' \in \mathcal{D}} \exp(\tilde{\beta}_{id't} - \alpha\sigma_{id't}^2)}.$$

- Beliefs update according to Bayesian Learning

$$\tilde{\beta}_{idt} = \tilde{\beta}_{idt-1} \frac{S_{it-1}}{S_{it}} + \bar{x}_{idt} \frac{f_{it}}{S_{it}} \quad (\text{Mean})$$

where  $\bar{x}_{idt} \sim \beta_d + \epsilon_{idt}$  and  $\epsilon_{idt} \sim N(0, \sigma_d^2 / f_{idt})$

$$\sigma_{idt}^2 = \frac{\sigma_d^2}{S_{it}} \quad (\text{Variance})$$

- Amount of learning for doctor  $i$  given by network structure and knowledge of all doctors

$$\begin{aligned} S_{it} &= S_{it-1} + f_{it} \\ &= S_{it-1} + I_{it} \times \sum_j \tau_{ij} S_{jt-1} \end{aligned}$$

- Doctors choose  $I_{it}$  to maximize PDV of expected utility

## Next Steps

- Assume  $I_{it} = 1$  for today. No endogenous investment in learning
- Only use Bayesian learning + discrete choice equations to estimate  $f_{it}$
- Then use these estimates of  $f_{it}$  with model of network learning to estimate  $\tau_{ij}$  and  $S_{i0}$  (and thus  $S_{it} \forall i, t$ )
  - Normalize  $S_{N0} = 1$
- Perform counterfactuals in simulated model about speed of learning w.r.t. different networks and initial knowledge
- Note: we have some theoretical results on endogenous learning and relaxing some homogeneity assumptions, but not with empirical results ready for today

## Bayesian Learning Estimating Equation

- Use  $S_{it+1} = S_{it} + f_{it+1}$  and define  $g_{it} := f_{it}/S_{it}$  to find

$$\begin{aligned}\tilde{\beta}_{idt+1} &= \tilde{\beta}_{idt} S_{it}/S_{it+1} + (\beta_d + \epsilon_{idt+1})f_{it+1}/S_{it+1} \\ &= \tilde{\beta}_{idt} (1 - g_{it+1}) + \beta_d g_{it+1} + \epsilon_{it+1} g_{it+1}\end{aligned}$$

- Substituting  $\tilde{\beta}_{idt} = \ln \pi_{idt} + \alpha \sigma_{it} + \eta_{it}$ , we find

$$\begin{aligned}\ln \pi_{idt+1} &= \ln \pi_{idt} \cdot \underbrace{(1 - g_{it+1})}_{=\rho_{it+1}} + \underbrace{\eta_{it}(1 - g_{it+1}) - \eta_{it+1}}_{=\kappa_{it}} \\ &\quad + \underbrace{\beta_d}_{=\delta_d} \cdot \underbrace{g_{it+1}}_{=\gamma_{it}} + \underbrace{\epsilon_{idt+1} g_{it+1}}_{=u_{idt+1}} \\ &= \ln \pi_{idt} \cdot \rho_{it+1} + \kappa_{it} + \delta_d \cdot \gamma_{it+1} + u_{idt+1}\end{aligned}$$

- Introduce reference drug  $d'$  and spells  $\tau$  to get main equation

$$\ln(\pi_{idt+1}/\pi_{id't+1}) = \ln(\pi_{idt}/\pi_{id't}) \cdot \rho_{it+1} + \delta_{d\tau} \cdot \gamma_{it+1} + u_{idt+1}$$

## Estimation results for persistence $\rho_{it}$ :

- Estimate by quarter; aggregate to molecule level
- Replace  $\delta_d \cdot \gamma_{it}$  with  $\delta_d + \gamma_i + \gamma_t$

Spell	1	2	3	4	5	6	7
Persistence—							
Mean	0.81	0.80	0.80	0.81	0.81	0.85	0.87
Std Dev	0.39	0.43	0.40	0.33	0.25	0.42	0.22
25 <sup>th</sup> pctile	0.69	0.68	0.68	0.70	0.72	0.71	0.78
75 <sup>th</sup> pctile	0.96	0.97	0.95	0.96	0.93	1.00	0.97
Doctor FE	Y	Y	Y	Y	Y	Y	Y
Drug FE	Y	Y	Y	Y	Y	Y	Y
Month FE	Y	Y	Y	Y	Y	Y	Y
Observations	787K	362K	558K	516K	1.4M	231K	4.3M

## Estimating $\tau_{ij}$ and $S_{i0}$

Combining network signal structure and Bayesian learning yields:

$$\begin{aligned}\hat{\rho}_{it} &= 1 - f_{it} / S_{it} \\ &= 1 - \frac{\kappa \times \sum_j \tau_{ij} S_{jt-1}}{S_{it}} \\ &= 1 - \frac{\kappa \times \sum_j \tau_{ij} \frac{S_{j0}}{\prod_{u=1}^{t-1} \hat{\rho}_{ju}}}{\frac{S_{i0}}{\prod_{u=1}^{t-1} \hat{\rho}_{iu}}},\end{aligned}$$

Parameterize  $\tau_{ij}(\boldsymbol{\alpha}) = \tau(\widetilde{dist}_{ij}, school_{ij}, cohort_{ij}, specialty_{ij}; \boldsymbol{\alpha})$  as follows,

$$\tau_{ij}(\boldsymbol{\alpha}) = \alpha_d \widetilde{dist}_{ij} + \alpha_s school_{ij} + \alpha_c cohort_{ij} + \alpha_{sp} specialty_{ij}$$

to get

$$\hat{\rho}_{it} = 1 - \frac{\sum_j \tau_{ij}(\boldsymbol{\alpha}) \frac{S_{j0}}{\prod_{u=1}^{t-1} \hat{\rho}_{ju}}}{\frac{S_{i0}}{\prod_{u=1}^{t-1} \hat{\rho}_{iu}}},$$

## Estimation results for initial knowledge $S_{i0}$ - preliminary

- Distribution of standardized  $S_{i0}$  highly skewed

Distribution of $\frac{S_{i0} - \bar{S}_0}{\sigma_{S_0}}$	
p1	-.110
p50	-.096
p75	-.067
p90	.025
p95	.155
p99	1.25
Skewness	24.5
Kurtosis	673

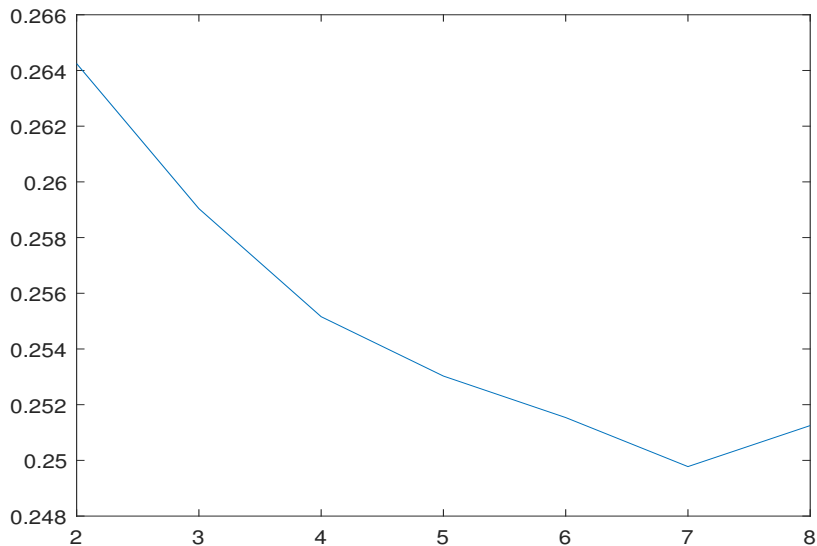


# Simulate the Economy

- Restrict analysis to first spell ( $D = 6$ ,  $T = 8$  quarters)
- Set  $\sigma^2 = \#$  (chosen to eyeball-match rate of convergence)
- Set initial period beliefs  $\tilde{\beta}_{i,d,1}$  consistent with initial period drug shares  $\pi_{i,d,1}$
- Set true  $\beta_d$  equal to that consistent with average of final period drug shares across doctors.
- Use estimated  $\alpha_d$  and  $S_0$  to construct  $f_{it} \forall i, t$
- Use Bayesian updating to construct  $\tilde{\beta}_{i,d,t} \forall i, d, (t > 1)$ 
  - Signals centered around truth, but finite signals means noisy updating
- Use  $\tilde{\beta}_{i,d,t}$  to construct  $\pi_{idt} \forall i, d, (t > 1)$

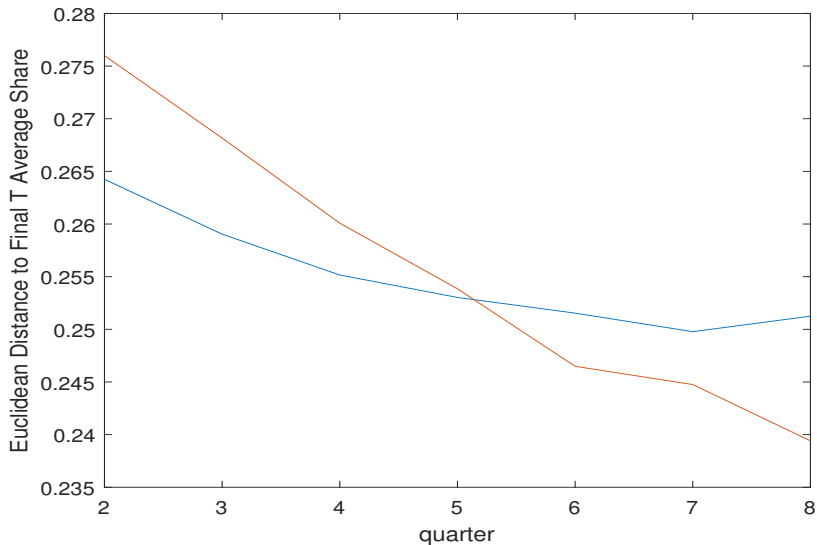
# Data: Drug Shares Converge Across Doctors

Euclidean Distance of Prescription Shares to Average  $T$ -shares



# Simulation: Drug Shares Converge Across Doctors

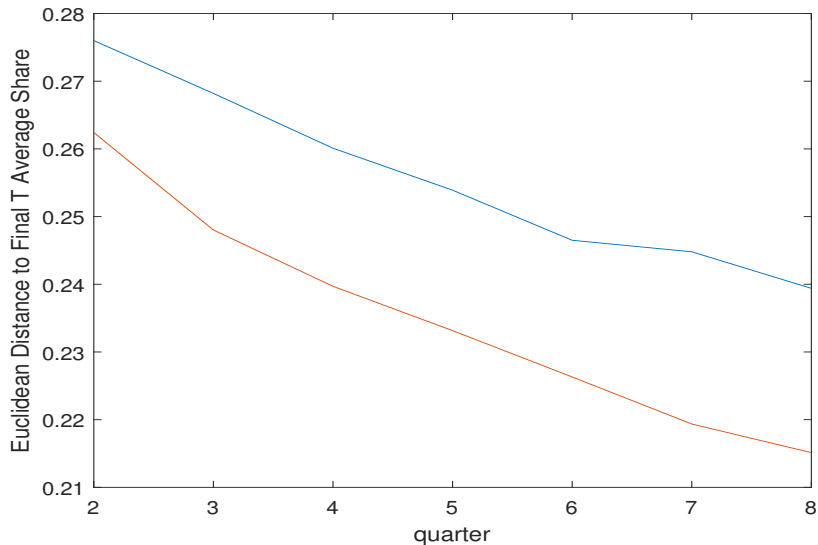
Euclidean Distance of Prescription Shares to Average  $T$ -shares



## Sim: Double Initial Knowledge Uniformly

Set  $S_{i0} = 2 \times S_{i0}^{\text{baseline}}$

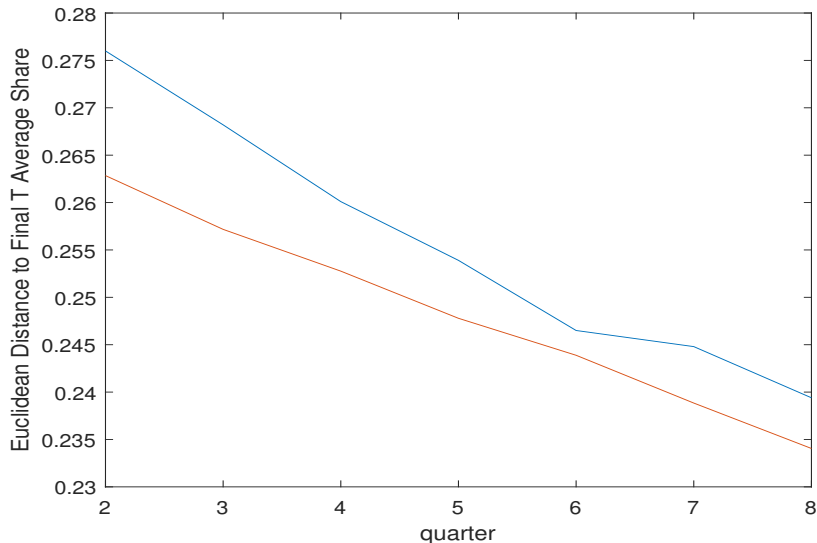
A bit less distance and more convergence. Meaningful only because not uniform network.



## Sim: Increase Initial Knowledge of Least Knowledgeable

Set all  $S_{i0} < p25(S_{i0}) = p25(S_{i0})$

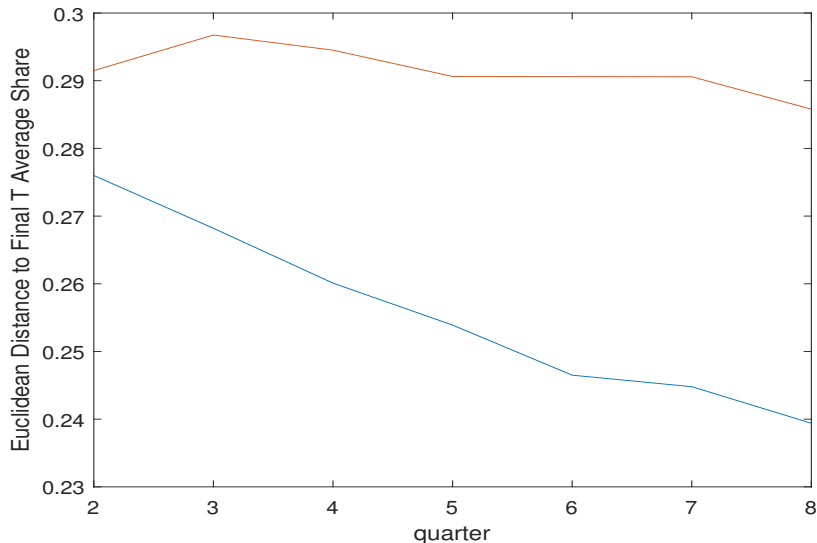
Less initial distance, but also less convergence.



## Sim: Decrease Initial Knowledge of Most Knowledgeable

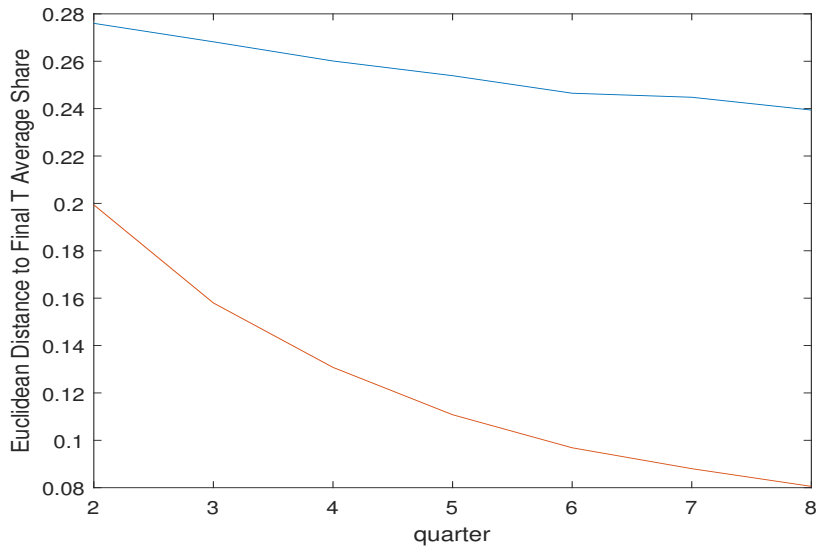
Set all  $S_{i0} > p75(S_{i0}) = p75(S_{i0})$

Much bigger distances, much slower convergence. Skewness.



# Sim: Drug Shares Converge Faster when Distance is Shorter

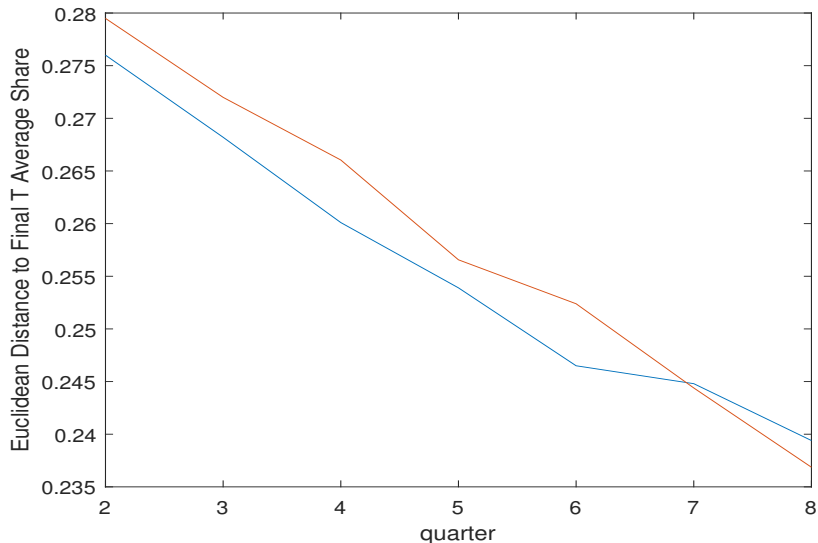
Set  $\tau_{ij} = 10 \times \tau_{ij}^{\text{baseline}}$



## Sim: Make Most Distant Doctors More Connected

Set all  $\tau_{ij} < p25(\tau_{ij}) = p25(\tau_{ij})$

Not much changes

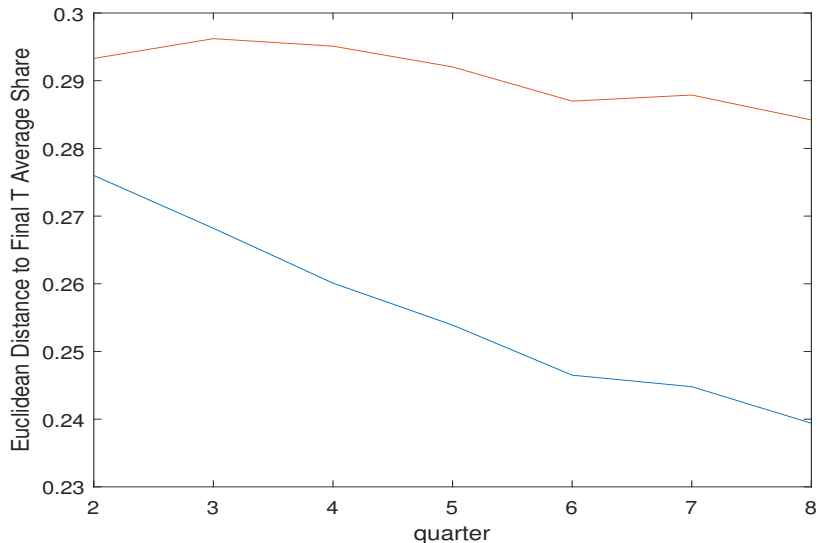




## Sim: Make Most Connected Doctors Less Connected

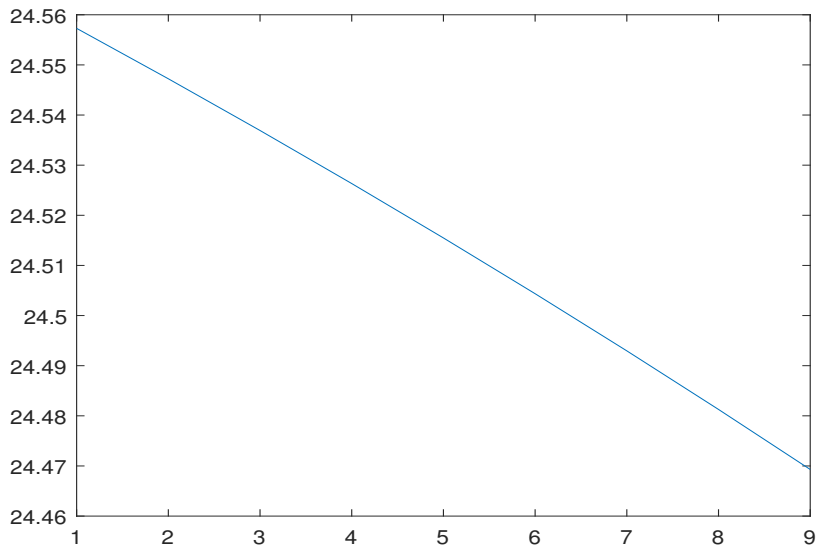
Set all  $\tau_{ij} > p75(\tau_{ij}) = p75(\tau_{ij})$

Much slower convergence



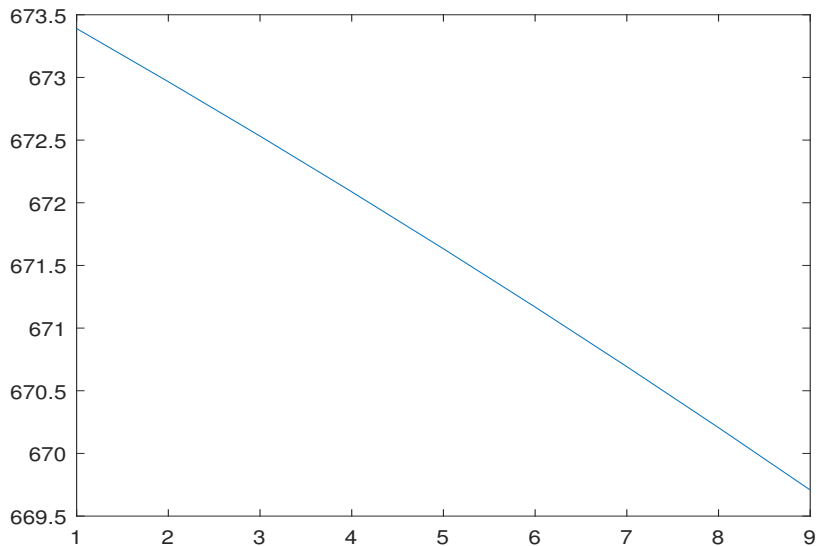
## Simulation: $S_{it}$ Skewness Decreasing Over Time

Diffusion of information is an equalizing force

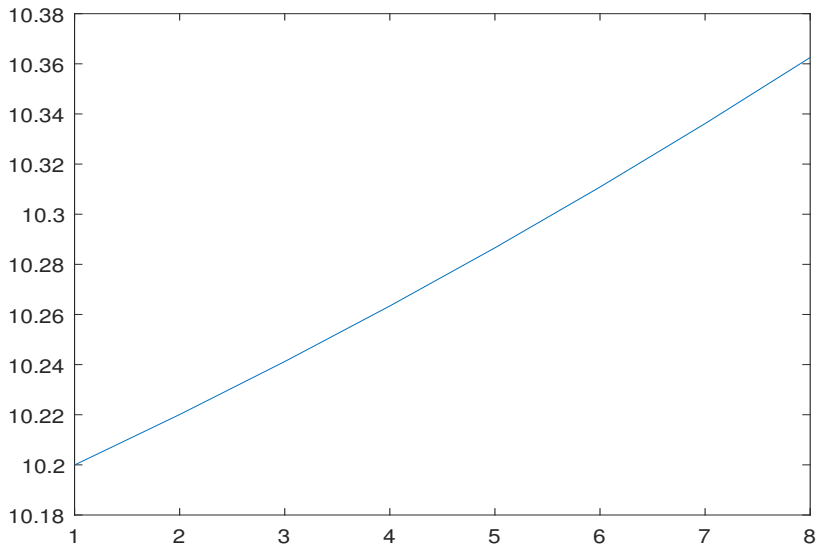


## Simulation: $S_{it}$ Kurtosis Decreasing Over Time

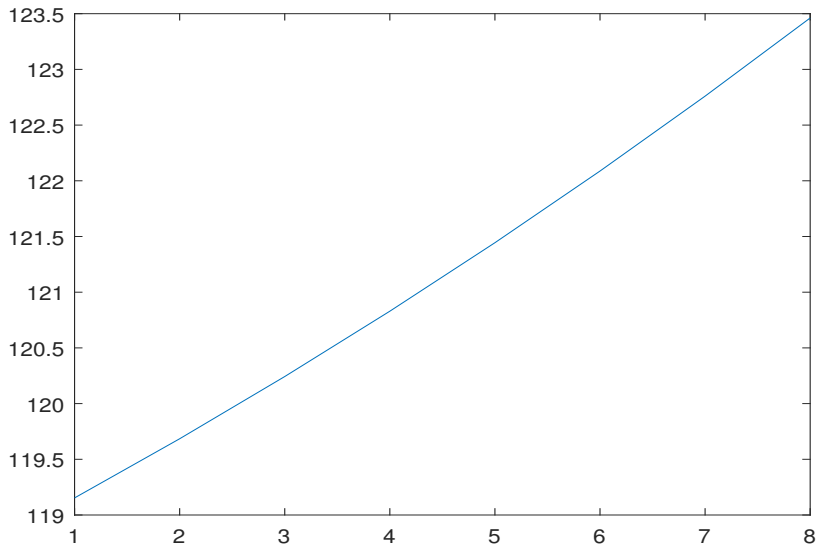
Diffusion of information is an equalizing force



## Simulation: $f_{it}$ Skewness Less Than $S_{it}$

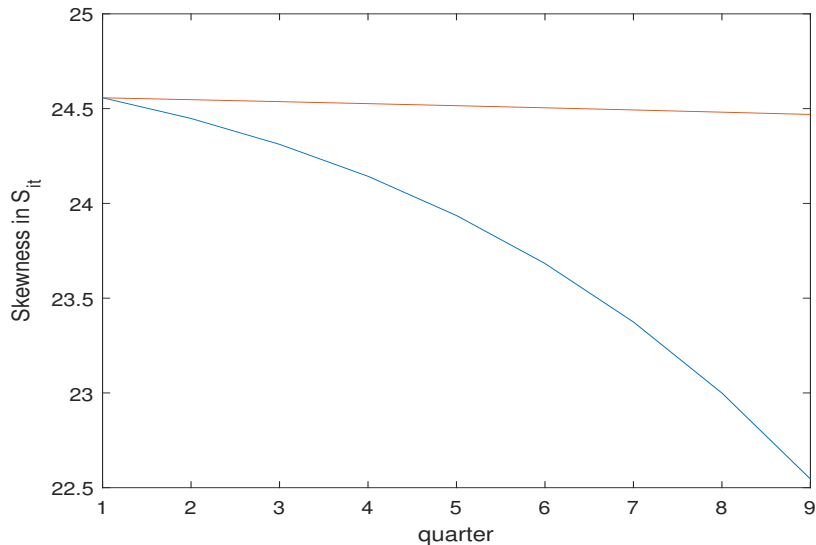


# Simulation: $f_{it}$ Kurtosis Less Than $S_{it}$



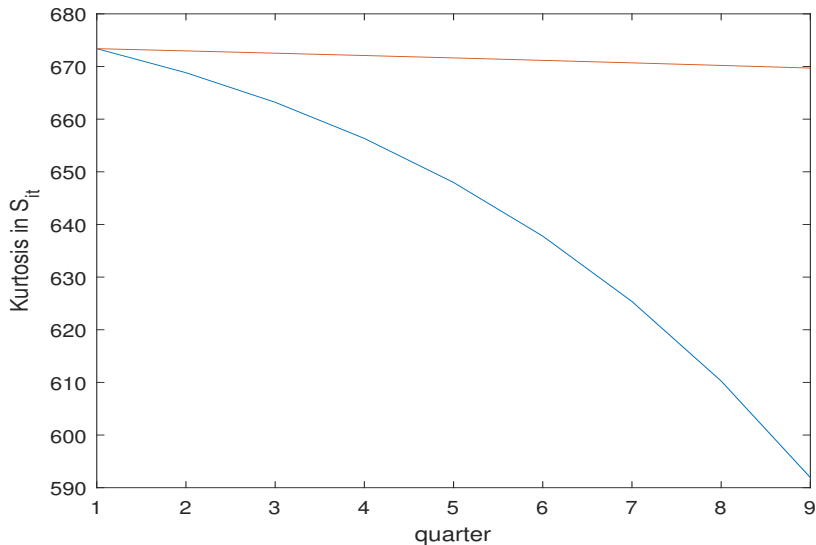
# Simulation: $S_{it}$ Skewness Decreases Faster When Distances are Shorter

Set  $\tau_{ij} = 10 \times \tau_{ij}^{\text{baseline}}$



# Simulation: $S_{it}$ Kurtosis Decreases Faster When Distances are Shorter

Set  $\tau_{ij} = 10 \times \tau_{ij}^{\text{baseline}}$



# Conclusion

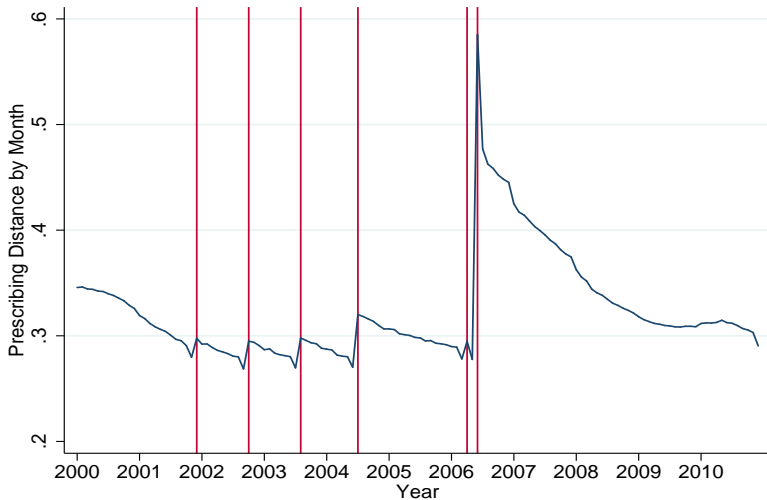
- Prescription shares converge over time, faster for central doctors
- Model-generated and estimated learning paths both indicate central doctors learn faster and that learning declines over time
- Many next steps:
  - Richer distance variables (cohort, medschool, specialty, facebook social connectedness of zipcode)
  - Learning from other doctors' beliefs
  - Risk aversion ( $S_{idt} \neq S_{id't}$ )
  - Endogenous investment in learning
  - Joint estimation of full model (not two step Bayesian learning and then network module), with more structural parameters ( $\beta_d$ ,  $\sigma_d^2$ , CARA, etc.)
  - Alternative models of the network
  - Introduction of new drugs—across spell analysis



# Appendix

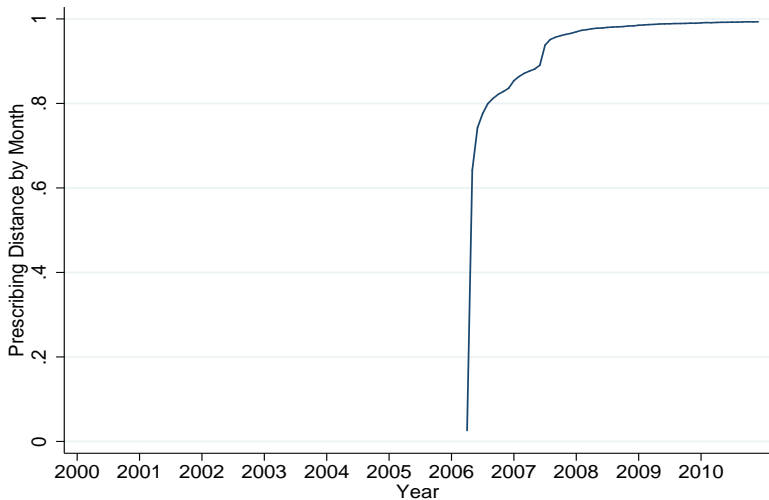
# Scripting Patterns Converge across Doctors within Zipcode

Convergence within zipcode suggests not just composition effect



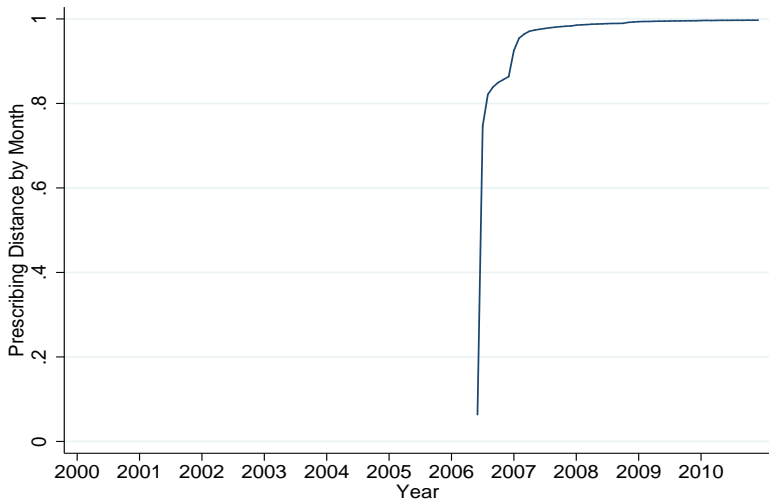
# Generic Prescription Share Converges

→ Within-molecule generic substitution, Pravastatin



# Generic Prescription Share Converges

→ Within-molecule generic substitution, Simvastatin



## Estimates of Average Convergence Rates

- Convergence target  $\bar{\pi}_{dT(t)}$  differs by spell,  $\bar{\pi}_{dzT}$  by zipcode-spell
- Control for prescribing intensity and doctor FE

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Dep variable:	Average Prescription Distance			
Drugs Target	All U.S. $\bar{\pi}_{dT}$	Six U.S. $\bar{\pi}_{dT}$	Six U.S. $\bar{\pi}_{dT}$	Six Zipcode $\bar{\pi}_{dzT}$
Time (months)	-0.0005 <sup>a</sup> (3.64e-06)	-0.0005 <sup>a</sup> (3.85e-06)	-0.0003 <sup>a</sup> (2.73e-06)	-0.0006 <sup>a</sup> (2.92e-06)
Rx Volume			-93.99 <sup>a</sup> (.0628)	-92.09 <sup>a</sup> (.0670)
Doctor FE	N	N	Y	Y
Observations	17334636	17334636	16241174	16241174

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Notes: *a* denotes 1% significance, *b* denotes 5% significance, *c* denotes 10% significance.

## Estimates of Convergence Rates: Generic Substitution

Dep variable:	Average Generic Share		
Molecule	Lovastatin	Pravastatin	Simvastatin
Target	1	1	1
Time (months)	.0013 <sup>a</sup> (1.79e-06)	.0051 <sup>a</sup> (5.12e-06)	.0038 <sup>a</sup> (3.68e-06)
Rx Volume	.5644 <sup>a</sup> (.1252)	15.05 <sup>a</sup> (.2340)	9.206 <sup>a</sup> (.1443)
Doctor FE	Y	Y	Y
Observations	7300484	4683682	6171908

Notes: *a* denotes 1% significance, *b* denotes 5% significance, *c* denotes 10% significance.

- Control for prescribing intensity and doctor FE