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# Improving diffusion in agriculture: an agent-based model to find the predictors for efficient early adopters

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## Abstract

Proven that the adoption rate of a new product is influenced by the network characteristics of the early adopters, the aim of this paper is to find the network features of the early adopters associated with high adoption rates of a specific new practice: the use of biodegradable mulching films containing soluble bio-based substances derived from municipal solid wastes. We simulated the diffusion process by means of an agent-based model calibrated on real-world data. Closeness and clusterization emerged as the most important network characteristics for early adopters to be successful. The results achieved represent the basis for the breaking down of a tailored diffusion strategy to overcome the psychological and socio-economic barriers of this kind of innovation within an environmental and sustainability-oriented transition policy in a rural context.

## Introduction

The bio-waste valorization is becoming an increasingly urgent priority for governments and environmental and social organizations (Morone et al. 2015). In this respect, the attention is mainly focused on the organic fraction of municipal and agricultural waste that can be used as raw material for biodegradable products (e.g., detergents, fuels, textile auxiliaries, plastic, fertilizers) (Motoneri et al. 2014, Scaringelli et al. 2016). Among these, a promising sustainable innovation is represented by biodegradable mulching films derived from soluble bio-based substances (SBOs) (Motoneri et al. 2014). This technology is at a development stage (TRL<sup>1</sup> 4/5), and its future adoption by farmers can improve both the sustainability of agricultural practices and the waste management process. This introduces the issue of innovation diffusion, proven that the achievement of the abovementioned benefits strictly depends on the spread of the novelty among a critical mass of users.

Here, we stress that, at the core of the diffusion process, there are operation and functioning of the social networks (i.e., the set and pattern of support, friendship, and communication relations) connecting people (Valente 1995). In fact, in its straightforward definition, the diffusion of innovation can be intended as “a special type of communication, in that the messages are concerned with new ideas” (Rogers 2003: 5). Communication implies the mobilization of social ties among people to create and share information and reach a reciprocal understanding (Rogers 2003). This process is fundamental in overcoming the innovation resistance that is the agents’ normal response against uncertainty and the costly readjustment activity imposed by the

innovation (Ram 1987). To put differently, the learning-from-others mechanism is likely to enhance the diffusion of innovation: this is true not only for innovations but also for farming strategies (Santeramo 2018).

More in-depth, social networks can influence diffusion providing (1) the medium for the circulation of *information* that is crucial to make the agents aware about the novelty and its real costs and benefits; (2) a certain level of redundancy of the information, deriving from *social reinforcement*, that is useful in overcoming uncertainty; (3) a certain level of *homophily* among actors (defined as the overlap degree of some agents attributes, such as education, socioeconomic status, and preferences) that favors common meanings, beliefs sharing, and mutual understandings (Rogers 2003).

As a consequence, the ability of the agents to affect their neighbors' (i.e., the others with whom they are connected) adoption decision is closely related to their role and location in the network. The social network analysis (SNA) has developed various centrality measures to capture network characteristics of the agents. These centrality measures can be considered by market operators and policymakers as a key driver to build effective promotional strategies. With this regard, the literature demonstrates that diffusion rates can vary greatly depending on the injection points (IPs) (i.e., the agents where the innovation is first injected, which are "early adopters") used, according to their network centrality (Banerjee et al. 2013). This is particularly true in the case of diffusion of new sustainable practices in agriculture (Tey and Brindal 2012).

In this perspective, this work represents a preliminary analysis of the network features which best predict effective IPs (i.e., those able to reach the highest adoption rates) in the case of a specific sustainable novelty, the SBO mulching films, in a rural context. The aim is to find those individual centrality measures that can be used as a rational criterion to select the best spreaders. The focus is on a community of farmers located in the north of Apulia (Italy) specialized in the production of vegetables. By means of an agent-based model (ABM), we simulated the effects of different IPs on the diffusion rates into an artificial community representing the one studied. In order to evaluate the robustness of our findings, we conduct a sensitivity analysis accounting for the possibility that the novelty disappoints a fraction of consumers. Moreover, we applied the SNA to obtain measures of IPs' network characteristics. Finally, we adopted a multiple linear regression model to estimate the effect of the various centrality measures of the IPs on the final adoption rates.

The use of social network potentialities to design successful diffusion campaigns is a topic largely investigated by the literature on ABMs of innovation diffusion (Goldenberg et al. 2001, Moldovan and Goldenberg 2004, Alkemade and Castaldi 2005, Goldenberg et al. 2007, Delre et al. 2007a). In this kind of approach, different from another kind of models such as linear programming, dynamic models, differential equation, the most elementary unit of modelization is represented by the single agent rather than the social system as a whole. This allows researchers to explicitly model the agents' heterogeneity, their social interactions, and their decision-making processes. In fact, the distinguishing feature of the ABM is that it represents the macro-level dynamics that take place in a social system as the result of the behavior of every single agent belonging to the social system and the interactions with its neighbors. For these reasons, ABM results are particularly suitable in showing the different performances of diffusion strategies based on the choice of the best connected actors (Valente and Davis

1999, Delre et al. 2007a, Goldenberg et al. 2009, Delre et al. 2010, Bohlmann et al. 2010, van Eck et al. 2011). Some of these models (Alkemande and Castaldi 2005, Goldenberg et al. 2000, Delre et al. 2007b; Bohlmann et al. 2010) explicitly include an adoption threshold to represent the different propensity of potential consumers of a new product to adopt (Kiesling 2012). Another type of models is devoted to simulating informational cascade (Banerjee 1992, Watts 2002). These models depict the graduation of the diffusion dynamic where different classes of consumer gradually adopt the novelty according to a progressive fashion distinguishing early adopters, followers, and laggards. When the novelty spreads in the largest part of the network, it occurs as a cascade.

The paper is structured as follows: the “[The diffusion model](#)” section describes the model employed, its basic assumption, its construction, and internal dynamics; the “[Simulation and results](#)” section reports the main results; and the “[Concluding remarks](#)” section concludes with some discussion and policy implications, including also suggestion for further research and some concluding remarks.

### **The diffusion model**

For the sake of clarity, in what follows, we shall describe the model according to the “guidelines for model development” outlined in Rand and Rust (2011) aimed at setting a rigorous procedure for AB modeling. The logical basis of the model is provided by the following assumptions that cover all the relevant elements of the model inferences; moreover, the references after each statement provide the empirical basis for the assumptions (East et al. 2016): (A1) Each agent has a specific innovation resistance, represented by an individual adoption threshold (Nisbet and Collins 1978). (A2) For the adoption to occur, the agent’s preference toward the innovation must overcome its innovation threshold. (A3) To form a preference, the agent must be aware of the innovation existence and its advantages (Chen 1996; Daberkow and McBride 2003). (A4) The agent grasps relevant information to become aware of the preferences of his neighbors (agents connected with it) (Molina-Morales and Martinez-Fernandez 2010; Narayan and Pritchett 1999; Van Rijn et al. 2012). (A5) The more homophilous the neighbor is, the more pieces of information the agent will grasp from it (Centola 2010, 2011). (A6) The higher the agent’s education, the higher its capacity to grasp information from its neighborhood as a whole (Gellynck et al. 2014; Tepic et al. 2012).

### **Model design**

The *scope* of the model is to reproduce the passing information dynamics and the adoption decision process among a population of farmers. In its most straightforward outline, the model depicts a network of *agents* connected by bidirectional *links*. As a consequence, the *environment* of each agent is represented by its neighbors that are the agents with whom it is connected via in/out-links through which it receives/sends information and influence. As highlighted by East et al. (2016), a common shortcoming in network diffusion models derives from the use of theoretical network structure (e.g., random networks, regular lattice) typically exhibiting the same centrality for all the actors. To overcome this limitation and to reach a better representation of relational complexity, the network in this case is modeled on the basis of a real-world network as explained below. The agents are divided into two classes, *ordinaries* and *IPs*; in each

model run, we have one IP, being the rest of the agent ordinaries. They are characterized by different *behaviors*: The ordinaries represent farmers that have to decide whether to adopt or not. The IP represents a farmer which has already adopted the novelty and uses it over all the model time span since the initialization phase. Concerning the set of *properties* of the model's elements, each link has a property representing the level of homophily between the agents connected [ $h$ ], and each agent has a preference toward the new technology [ $p$ ], an adoption threshold [ $\theta$ ], a level of education [ $e$ ], and the innovation status [*novelty?*]. These properties represent also the basic *inputs* of the model, while the model fundamental *output* is the number of ordinaries that have adopted at each time step.

### Model construction

The model is structured into two phases: the initialization and the iteration. At the *initialization* step,  $n$  agents and  $m$  links are created. At this phase,  $p$  is set at 0 for ordinaries and 1 for IPs and  $\theta$  and  $e$  are set at specific values (see below). The innovation status is set *false* for ordinaries and *true* for the IP. At each *iteration* time step, each ordinary (1) reconsiders its preference after having received information from its neighbors,<sup>2</sup> (2) pass information to neighbors,<sup>3</sup> and (3) decides to adopt (i.e., *novelty?* is set *true*) if  $p \geq \theta$ . This sequence is repeated until the model time span reaches the value  $t$ .

Concerning step 1, it is worth noting that the sole factor influencing the preference of an ordinary agent is the preferences of its neighbors. Specifically, at each time  $t$ , each ordinary  $i$  calculates its preference  $p_{it}$  as the sum of its preference in the previous period  $p_{it-1}$  and the average of the preferences of its  $j$  neighbors  $p_{jt}$  weighted with the homophily degree with its neighbors  $h_{ji}$ . This sum is then corrected multiplying it by the level of education of  $i$   $e_i$ . Formally, the calculation of  $p_{it}$  is:

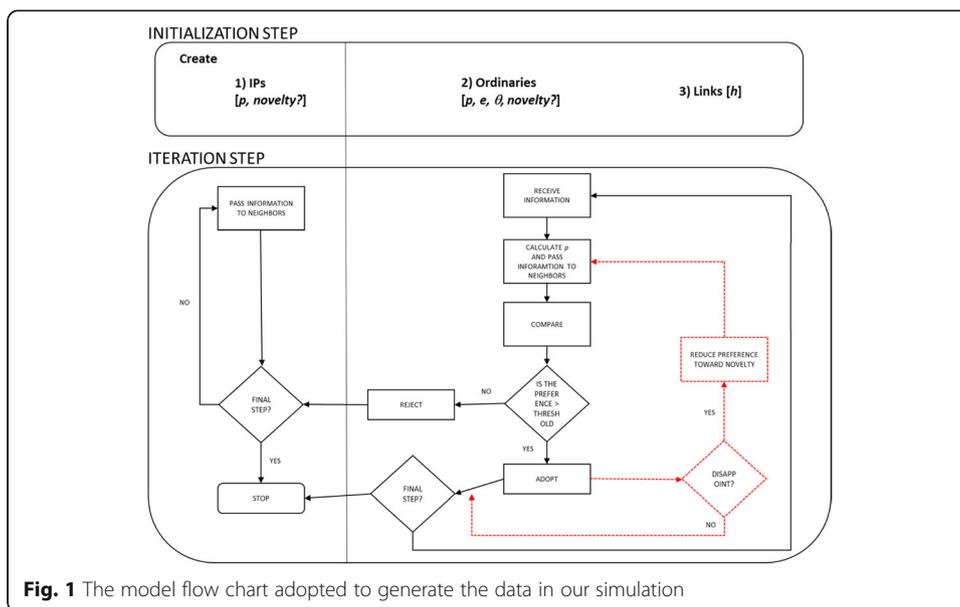
$$p_{it} = p_{it-1} + \left( \sum \frac{p_{jt-1} h_{ji}}{n} \right) C \text{ with } C = \frac{e_i}{\max_e}. \quad (1)$$

It is worth noting that the time dimension in this process acts as an accumulator of the preference level allowing to unravel the interaction between network features and preference formation.

To augment the model reality, we developed also a disappointment extension, which introduces a common issue in the innovation adoption process, that is the probability that the novelty disappoints a fraction of users. To simulate this behavior, the disappointment version includes the possibility that the innovation disappoints the adopter. When this case occurs, the model provides a possible third action for the ordinary agent, to reduce its preference of a variable % (between 25 and 75%) toward the novelty. In this version of the model, the disappointment is modeled as a random event which affects a variable fraction of the population (between 0 and 25%).

### Model verification

Our model was implemented and run using the NetLogo 5.2 platform (Wilensky 1999). To import the network under investigation, we included in the model a routine adapted by the "Network Import Example" authored by Uri Wilensky and available at the modeling commons platform.<sup>4</sup> The sequence of model execution is outlined in Fig. 1. It plots the chart flow of the model dynamics that is the internal agent's decision logic.



**Fig. 1** The model flow chart adopted to generate the data in our simulation

The code flow was analyzed, and each piece of the code was tested to verify its correct functioning. In this phase, we also employed corner cases (extreme values of the inputs) to verify that the implemented model does not show aberrant behaviors (e.g., no adoption is realized without any IP).

**Model calibration**

The identification of model parameters was based on a case from South Italy, located in the Province of Foggia. It relates to a group of specialized vegetable farmers ( $N = 80$ ) which already use conventional or biodegradable mulching technique and are therefore potentially interested in the innovative use of SBO-derived films (Scaringelli et al. 2016). Sample size covers 2% of the population. This is a reasonable sample extension to obtain enough real-world data for the calibration of the model parameters. This case study is suitable to this end, prove that it offers a cross-section of the social and professional interactions of vegetable crop farmers in this territory. Being the technology under investigation still at a development stage, no real adoption rate already exists. This is exactly what the ABM model addresses: to simulate the effect of various IPs on the final adoption rate. Thus, we used the case study with the sole purpose to calibrate the following model parameters: (i) the network structure, which is defined by the number of agents, the number of links connecting the agents, and the links distribution among agents; (ii) the agent parameters  $e$  and  $\theta$ ; and (iii) the link parameter  $h$ . We obtained the information on the network structure by means of deep face-to-face interviews with two experts which live and work within the context of the case study and can be considered direct observers. These interviews were directed at recognizing the social and professional relations connecting the web by means of a participatory social network observation.<sup>5</sup> To this end, the experts were asked to trace the who-knows-who relations and to identify the affiliations to local cooperatives. To obtain the other information needed to calibrate the reminder of model parameters, we performed a questionnaire survey on the actors forming the network asking for age, years of education,

cropping patterns, willingness to adopt SBO mulching films, farm size, and number of employees. We also calculated the distance between the farmers based on their location. Some descriptive statistics are reported in Table 1.

The relational data were used to build the network of firms that form the interaction arrangement of the agents. This network is formed of a single component, not fragmented, with a medium level of density (21%). Within this network, two randomly chosen nodes have an average distance of 2.42, and the maximum distance observed is 6. On the whole, the network results are very clustered (clustering coefficient 0.78). The demographic data were employed in the validation of agents' properties. Specifically, we used the following parameters:  $n = 80$ , as the number of respondent farmers. In each simulation, we split this number in 79 ordinaries and 1 IP;  $m = 1296$  that is the number of links observed in the real farmers' network. Moreover, in the construction of the simulated network, the links have not been evenly distributed among the agents. Instead, the agents in the model reproduce exactly the relational profile of each farmer interviewed, with the same number of relations with the same neighbors;  $h$  varies in the range [0.1]. Rather than using a single dimension, we set this property on the base of four socioeconomic attributes (farm size, number of employees, age, and distances between farmers), as in Blau et al. 1984, Mc Pherson et al. 2001; Centola et al. 2007;  $p$  is 0 for ordinaries and 1 for IPs; *novelty?* is *false* for ordinaries and *true* for IPs;  $\theta$  reflects the willingness to adopt SBO mulching films declared by respondents and is set on a six-degree scale ranging between  $-1$  (completely adverse to adoption) and  $1$  (completely favorable to adoption);  $e$  is various and reflects the year of education of each farmer. For the estimation of  $t$  (i.e., the model time span), we matched a 5-year time horizon (representing a rational span to allow the innovation to become mature) with 45 model periods, that is the time needed by the most part of the IPs to reach a stable adoption rate. As a consequence, in our model, a time step is equal to 40 days.

## Simulation and results

The procedure adopted to find those centrality measures that can be used as rational criterion to select effective IPs can be broken down in three main phases: (1) calculation of the centrality measures for each agent in the network, (2) simulation of the IP's effect in terms of final adoption rate using the ABM model, and (3) estimation of the effect of centrality measures on the simulated adoption rates employing the multiple linear regression.

### Phase 1

By means of the SNA, we calculated six network measures for each agent (Table 2).

**Table 1** The descriptive statics of observations of our case study

	Farm size (ha)	Employees	Age (years)	Distances between farmers (km)	Education (years)
Mean	76.96	13.94	45.74	59.77	10.84
Standard deviation	203.71	16.76	11.6	40.92	3.53
Min	4	1	24	0	5
Max	1805	112	72	198.42	18

Source: our elaboration

**Table 2** The network measures employed to characterize the IPs

Network measure	Description
1) Degree centrality	It expresses the number of links of a given node. It directly affects the capacity of a node to immediately influence the information flow in the network (Wasserman and Faust 1994).
2) Betweenness	It measures how often a given node is between two other nodes along their shortest path, that is how many times a node acts as a bridge. This measure can be interpreted as an indicator of how the single node plays the role of "gatekeeping" (Borgatti et al. 2013).
3) Closeness	It is the reciprocal of the farness of a node. It is expressed formally as the inverse of the sum of the distances from a certain node to all others in the network. The measure here used is a normalized version where the reciprocal of farness is expressed as a percentage of the minimum possible farness (Wasserman and Faust 1994).
4) Average reciprocal distance (ARD)	It measures how the node is close to the whole network. For a given node, it is calculated as the summation of all other nodes weighted for the reciprocal of the farness from the given node. The greater the value of this indicator, the greater is the "connectedness" of the node (Borgatti et al. 2013).
5) Local clustering coefficient	It is the density of the neighborhood of the given node. It ranges between 0 (any neighbor is connected with another) and 1 (the neighbors are fully connected) (Newman 2003).
6) Eigenvector	It looks at how close is the node with respect to the whole network, putting less importance on more "local" closeness. It is calculated by means of a factor that identifies dimensions of the distances among actors. The first dimension identifies the "global" aspects of distances among actors, while the other dimensions capture more specific features (Hanneman and Riddle 2005).

Source: our elaboration

Table 3 contains the descriptive analysis of the abovementioned network measures. All the measures used were standardized over a range of 0.1 to facilitate the interpretation of the results. The table shows that each node (1) is connected to 21% of the rest of the network in mean, (2) intercepts 2% of the shortest path length among others, (3) is rather close to others, (4) has a medium level of ARD, (5) is characterized by a high local clustering coefficient, (6) and has a relatively low eigenvector.

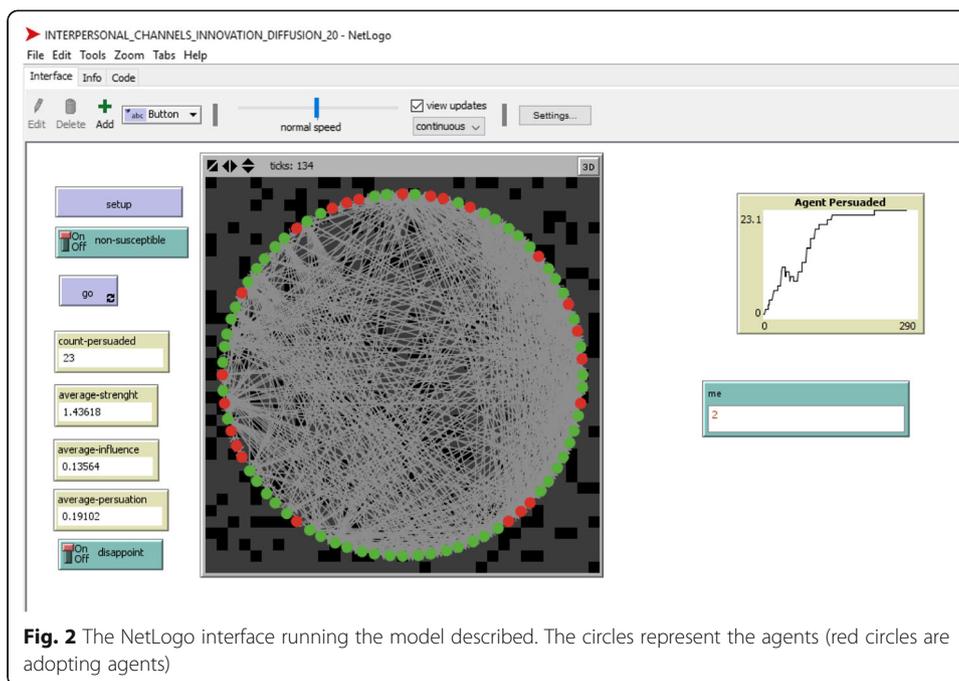
## Phase 2

We used the NetLogo platform to code and run the ABM model. Figure 2 depicts the interface of the model implemented. In the simulation, each one of the 80 agents is used as IP alternatively; moreover, the robustness of the results is proved simulating the diffusion process repeatedly using each IP many times (batches of 20 runs) and taking the means of the simulated data.

**Table 3** The table synthesizes the descriptive statistics of the SNA network variables

Network measure	Abbreviation	Mean	St.dev	Min	Max
1) Degree centrality	DEGREE	0.21	0.13	0.01	0.57
2) Average reciprocal distance (ARD)	ARD	0.40	0.09	0.00	0.54
3) Eigenvector	EIGENV.	0.13	0.09	0.00	0.33
4) Betweenness	BETWEEN.	0.02	0.03	0.00	0.10
5) Local clustering coefficient	CLUST.COEF.	0.76	0.27	0.00	1.00
6) Closeness	CLOSEN.	0.43	0.09	0.22	0.60

Source: our elaboration



Since the main output of the simulation is the number of adopters at the final time, we simulated the number and rate of adopters for each IP at  $t = 45$  (Table 4) for both versions of the model (basic and disappointment).

We also performed a robustness check of results, simulating the number of adopters and calculating the adoption rates also at  $t = 68$  and  $90$ . The variation is negligible. This means that at  $t = 45$ , the process is at its climax with a little part of the dynamic already working. We used the adoption rate at  $t = 68$  and  $90$  in the estimation of centrality effect (see the “Phase 3” section below) with no variation in the sign and significativeness of coefficients and with almost the same values. We verified also that the threshold at  $t = 45$  is the best end for the model simulation proven that the adoption rate at this time step displays a little more variability than  $t = 68$  and  $90$  magnifying the difference in the variables estimations.

The number of adopters is around 19 for the base model and 11 for the disappointment version. For both versions, the level of dispersion is very low, that is, excepting little variation, the IPs work in a rather same manner. The maximum rate of adoption is 25% for the basic model and 17% for the disappointment model.

At least for this case study, these results indicate that the individual characteristics of the IPs are not primarily responsible for the diffusion process. Instead, the density and clusterization of the network do the most part of the diffusion work.

**Table 4** The table shows the number and rate of adopters at the final step of our simulation

Model specification	Mean (%)	St.dev (%)	Min (%)	Max (%)
Base	19.40 (0.23)	2.34 (0.03)	9.30 (0.10)	20.90 (0.25)
Disappointment	10.85 (0.12)	1.57 (0.02)	5.70 (0.06)	14.30 (0.17)

Source: our elaboration

### Phase 3

In this phase, we estimated the effect of the centrality measures calculated in phase 1 on the adoption rates (AR) simulated in phase 2, by means of a simple model of linear regression, specified as follows:

$$AR = \alpha + \beta_1 \text{Centrality} + \beta_2 \text{ARD} + \beta_3 \text{Eigenvector} + \beta_4 \text{Betweenness} + \beta_5 \text{LCC} + \beta_6 \text{Closeness} + \varepsilon$$

where AR is the adoption rate in the base and disappointment models (cfr. Table 4), and the regressors are those specified in Table 3, with ARD standing for average reciprocal distance and LCC standing for local clustering coefficients.

Table 5 reports the results of the regression. What we can firstly observe is that the positive influence of a dense network on the innovation diffusion is confirmed by a high intercept in the base model. The analysis of this model also highlights degree and closeness centralities as the best indicators for effective IPs. Both these measures have in fact a positive effect on the adoption rate and are highly significant. Moreover, their impact is remarkable; an augment of 1% of degree and closeness centrality produces an increase of 30% and 47%, respectively, in the adoption rate. On the contrary, betweenness and eigenvector centralities exhibit a negative. This is a surprising result, proven that betweenness is a measure of how many times an agent acts as a bridge and is considered an important indicator of how the single agent plays the role of broker, while eigenvector is a measure of how central the neighbors of the agent are. High levels of these centralities are supposed to reflect into a great influence on the network with a potential crucial role in passing information and innovation spread dynamics (Borgatti et al. 2013; Hanneman and Riddle 2005). Finally, despite the average reciprocal distance and the clustering coefficient in particular are important network measure capable of providing social reinforce in the new behavior diffusion (Centola 2011) in this case resulted not statistically significant.

These partly counterintuitive results are probably due to the specific structure of the real network analyzed. Most part of the literature, indeed, analyzed the effectiveness of the centrality measures on a completely theoretical network (i.e., regular lattice, random, small world), but the morphology of the network is clearly responsible for the

**Table 5** The table synthesizes the results of the econometric analysis

Variable	Base		Disappointment	
	Coef.	Std.Err.	Coef.	Std.Err.
CONST.	11.397***	2.6827	2.2744	2.1445
DEGREE	30.028***	10.7661	0.1068	8.6063
ARD	-29.972	28.973	26.4463	23.1607
EIGENV.	-46.4384***	9.0967	-14.9239**	7.2718
BETWEEN.	-49.2363***	13.5898	-20.2023*	10.8635
CLUST.COEFF.	0.6331	0.9955	0.8556	0.7958
CLOSEN.	47.4929**	22.5804	-1.497	18.0505
R <sup>2</sup>	0.3504		R <sup>2</sup>	0.2445
Observations	80		Observations	80

\*Significant at the 90% level

\*\*Significant at the 95% level

\*\*\*Significant at the 99% level

whole diffusion dynamic and of the actual power of the single positions. In the network under investigation, the agents with the highest betweenness act as bridges between two major components of the network. These two components differs in size and structure, being the greater (60 nodes) 3.75 times the size of the smaller (16 nodes). Moreover, the former has a pretty random structure highly centralized with many long ties, while the latter is a very clustered network with a highly regular structure. In this context, the most part of the diffusion dynamics happens in the greater component, making relatively more important role of central and close IPs in its core than the role of brokers with the minor component. The supremacy of the network structure over the diffusion dynamics is also confirmed by the results from the disappointment model: when harsher conditions to innovation adoption emerge with disappointment possibility, the positive influence of IPs centrality on adoption rates ceases, while the negative operation of betweenness and eigenvector remains, though reduced.

Being the model we used for the estimation a multiple regression, the findings related to each single centrality measure should not be intended as separate and self-contained but as parts of a comprehensive frame capable of leading to a multifaceted and clear profile of the ideal spreader: central and close to the rest of the network but not “between” the most part of actors, with no or limited bridge toward other very central actors. In other words, in this specific structure, the diffusion of innovation is accelerated when the spreader can exert direct influence through immediate links on a large part of the network and is hindered when this influence is mediated by others.

### **Concluding remarks**

This paper aimed at performing a preliminary analysis of the network features characterizing IPs capable of reaching high adoption rates in the case of the adoption of SBO mulching films in a rural context. To this end, we combined techniques from SNA, ABM, and linear regression to calculate the centrality measures of the spreaders, simulate the adoption rates of the innovation, and estimate the effect of centralities on the adoption dynamics. These results should be viewed as a first step in identifying the network measures featuring good IPs. We found the most straightforward centrality measures (i.e., centrality degree, closeness) as the most relevant in spreading diffusion. This is not a trivial result proven that more “insightful” measures (i.e., betweenness and eigenvector) resulted to exert an opposite effect on the adoption rates. This contributes to addressing the initial problem of finding those individual centrality measures suitable to identify operative criteria to select the best spreaders. Rather than single specific criterion, we identified an actual profile of the best spreader.

We found that, for the diffusion of sustainable innovation in agriculture, and at least in the case of context and relational structure similar to that studied, it is primarily important how connected the IP is (how high is its immediate influence) and how relatively close it is to the rest of the web. Another major point is the emerged supremacy of the network structure on the diffusion dynamics. It makes the difference on the effectiveness of IPs positions creating positive or negative chance to innovation diffusion. This is very important for policy and marketing decisions, especially in rural areas which are characterized by a highly specific social structure that should carefully be considered to design a successful spreading campaign. The implications of our paper are relevant in terms of rural planning and development. To the extent that innovations are potential drivers of competitiveness,

policy interventions should be planned in order to achieve not only efficacy, but also efficiency, exploiting the leverage effect of social capital (Nardone et al. 2010): targeting the best early adopters, capable of enhancing the learning-from-others mechanisms (Santeramo 2018) would help in lowering the costs of policy interventions and increasing the impacts of the rural development measures.

Finally, the fact that no agent reached at least a 25% adoption rate highlights the need to identifying the optimal number of injection point (rather than a single injection point) to maximize the potential diffusion of the technology, also considering the fact that it is costly for advisors to spread the technology among end users. The kind of model used in this paper might help in the design of balanced groups of IPs. A caveat of this study is represented by a limited case study with specific features (density and clusterization). The effects of the individual characteristics on the final rate of adoption should be interpreted on this basis. A more comprehensive analysis should include the investigation of different network structures (e.g., high vs. low density, regular vs. randomized structure, high vs. low average degree). Another area of analysis is represented by the measurement of the impact of exposure number per agent to the word of mouth. The expected result of such a deeper investigation is the achievement of valuable hints for marketing and policy-making actors in the light of the bio-waste valorization.

## Endnotes

<sup>1</sup>TRL is the acronym of “technology readiness levels” that is a system to assess the maturity level of a specific technology developed by the National Aeronautics and Space Administration (NASA) see [https://www.nasa.gov/directorates/heo/scan/engineering/technology/txt\\_accordion1.html](https://www.nasa.gov/directorates/heo/scan/engineering/technology/txt_accordion1.html)

<sup>2</sup>Specifically, each ordinary recalculates its  $p$  as the sum of its  $p$  in the previous period and the average of the  $p$  of its neighbors weighted with the homophily degree ( $h$ ). This sum is then corrected multiplying it by the level of education ( $e$ ).

<sup>3</sup>This kind of action is the only one that applies to IPs.

<sup>4</sup>The Network Import Example is downloadable at [http://modelingcommons.org/browse/one\\_model/2214#model\\_tabs\\_browse\\_procedures](http://modelingcommons.org/browse/one_model/2214#model_tabs_browse_procedures)

<sup>5</sup>This method is based on the involvement of actors directly implicated in the network investigated, by means of workshops or deep interviews to co-produce a representation of that network (Edwards et al. 2010).

## Acknowledgements

We gratefully acknowledge the comments provided by G. Giannoccaro on a previous version of the present research.

## Funding

We have not received funding for thi researchers.

## Availability of data and materials

Data are available from the authors upon request.

## Authors' contributions

AL has coordinated the work and is responsible for the “Introduction,” “The diffusion model,” and “Simulation and results” sections. FS is responsible for the “Concluding remarks” section. AB has provided excellent research support. All authors read and approved the final manuscript.

## Competing interests

The authors declare that they have no competing interests.

## Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Received: 24 November 2017 Accepted: 10 January 2019

Published online: 28 January 2019

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