

Individualized Cancer Therapy, what is the Next Generation?

Da-Yong Lu^{1*}, Ting-Ren Lu¹, Jin-Yu Che¹ and Nagendra Sastry Yarla²

¹Shanghai University, Shanghai, PRC, China

²Divisions of Biochemistry and Chemistry, City University of New York School of Medicine, Convent Avenue, New York, USA

*Corresponding Author: Da-Yong Lu, Shanghai University, Shanghai, PRC, China.

Received: April 08, 2018; Published: June 26, 2018

Abstract

Cancer is the secondary most frequent cause of disease-induced human mortality worldwide. Yet, it is different disease with a universal feature of unlimited growth, sometimes invasive and remote metastasis (> 100 different pathogenic types). Individualized cancer therapies/personalized cancer therapies (ICT/PCT) are targeted to cope with the tricky character of disease diversity. The next generation of ICT/PCT must simultaneously contain information of pharmacology (drug sensitivity), oncology (tumor etiology/pathological information), computational network and patient's desire (decision-aid). Any unilateral efforts could not be able to reach maximum beneficial outcomes in clinics. This future trend is inevitable. In addition, anticancer drug combination might undergo great changes and optimum. After all, integrated clinical ICT/PCT systems (collective personalized paradigms) might make a difference for cancer treatments in future.

Keywords: Individualized Cancer Therapy; Personalized Medicine; Pharmacology, Neoplasm Pathology, Drug Combinations; Neoplasm Metastasis; Decision Aid

Backgrounds

Cancer as the secondary most frequent cause of disease-induced mortality causes annual human mortality of 7 - 10 million world-wide [1,2]. As cancer is a different disease, the hallmarks of cancer pathogenic traits vary in tumor genomic mutation/translocation/abnormality and with tissue origin/disease stages in the clinic [3,4]. Several types of individualized cancer therapy/personalized cancer therapy (ICT/PCT) have been designed for choosing anticancer drugs in the clinic [5-10] - include drug sensitivity test (DST) [11-13], tumor biomarkers/bioinformatics detection [14-18], pharmacogenetics (PG) [19-21], individualized antimetastatic therapy [22,23], drug combinations [24], cancer assistant therapy [25,26], patient's decision aids [27,28] and so on [29-31]. Some of them, such as DST have been established over 60 years.

Major characters of current ICT/PCT disciplinary

Therapeutic benefits vs survival benefits by DST utility in the clinic

From the surface, drug responses (partial response-PR or complete response-CR) in cancer patients are improved by DST utility. However, only less than 25 - 30% clinical therapeutic data state a survival benefits by current ICT/PCT. In most cases, patients' survival is almost the same in spite of DST or other ICT/PCT discipline utilities [11-13]. In addition, these kinds of survival elongations are only several weeks/months that is far beyond our requirement-long term disease-free for late-stage of cancer patients.

Biomarkers and bioinformatics

Bioinformatics is a modern approach that provides by a variety of techniques (omics) for mining and analyzing overall tumor abnormalities of DNA, RNA, proteins and glycoligands in human bodies [14-18]. Presently, the best example of utilizing cancer biomarkers or bioinformatics is for revealing abnormal proteins of cancer antigens. Though previous practice has established workable routines (more than 10 biomarkers) in general hospitals, it is not perfect for drug choice.

Why do we give this conclusion? If we know all information of tumor origin and pathological progresses, we still cannot determine which type of anticancer drug is most suitable. Anticancer drug activity is not parallel with oncology in mathematical ways. The pharmacological data of drug is more complicated comparing with mathematical/physics data analysis of oncologic changes.

Pharmacogenetics (PG) for cancer therapy

By entering this millennium, clinical applications of PG have been greatly intensified worldwide, especially PG for hepatic or other organ metabolism enzymes in human bodies. Presently, the purpose of PG anticancer study is mostly to predict the fraction of active or inactive metabolites. The final required dosage of a drug is deduced from the genetic variation of metabolic enzymes in tumors or human bodies [19-21] (Figure 1). Overall, PG study at this stage is an effort to maximize efficacy and minimize toxicity of drugs in individual patients. Yet, present PG can only detect a narrow-range of human genes and improperly judgment of anticancer activity of drugs in most patients.

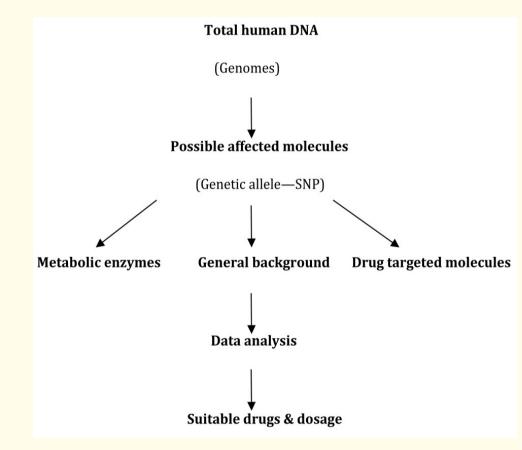


Figure 1: The schematic diagrams of cancer therapeutic PG.

Individualized antimetastatic chemotherapy

Similarly, large portion of cancer mortality (90%) is caused by cancer metastasis [32-35]. However, DST commonly evaluates drug response to primary tumor. No specific therapy against metastatic tumors fails for improving patients' survival. This shortcoming will greatly undermine current ICT/PCT imperfection.

Since cancer metastasis plays key role for cancer deaths, it is reasonable to garner growing attentions of neoplasm metastasis from individual patients. Individualized antimetastatic chemotherapy is to treat cancer according to different stages of neoplasm metastasis [22,23].

However, until now, clinical cancer treatments are mainly focused on primary tumors. Antimetastatic drugs are not widely developed [36-43]. Due to this knowledge deficiency, any small breakthrough for antimetastatic therapy is supposed to make a great difference in cancer therapies. Apart from this antimetastatic drug development, antimetastatic drug targets [44-61], especially against formed metastatic foci are top priorities in cancer treatment study. Some new PCT/ICT modality with this function can be valuable avenues for this breakthrough.

Therapeutic dilemma in metastatic treatment has been noticed. More recently, it is known that transmission of primary tumor into mobile status (floating in the vessels) is the transmission from epithelial to mesenchymal (EMT) and transmission of mobile tumors to metastatic nodules in remote organs is from mesenchymal to epithelial (MET) [62-66]. Thus, it might be mechanistically therapeutic oppositions between primary tumor and formed metastatic tumors.

Primary tumors → EMT → Circulating tumors → MET → Metastatic nodules

Figure 2: Overall picture of pathological mechanisms from primary tumors and metastatic tumors [74].

There is an opposite biological pathways and mechanisms between primary tumor and metastatic tumors. So, it is proposed that anticancer agents inhibiting primary tumors might be a promoter to metastatic tissues [67,68]. Facing this therapeutic dilemma, we argue that drug combinations/biotherapy may be helpful for this clinical situation [68-71].

Drug combinations

Most cancers have multiple genetic alterations and molecular abnormalities, especially for late-stage cancer patients. Drug combination might help us on this matter [72,73]. Generally speaking, anticancer drug cocktail might be one of the good solutions for cancer therapy [74-78]. However, it becomes a modern cliché in clinical cancer treatments [74-78]. How to combine use of anticancer drugs is an emerging problem for anticancer drug therapy study [79,80]. Future trends will be discussed in later section.

Assistant cancer therapy

Similarly, anticancer assistant therapy is also a useful way of therapeutic improvements, especially for some very refractory cancer types-solid tumors [25,26,55-61]. In future, some new laws (what types of anticancer or antimetastatic drugs are the best combinations) remain to be elucidated. These types of medical findings must be repeatable and discovered by experimentally pharmacologic study and double-blind clinical evaluations [25,26]. These topics are the major pharmacological/therapeutic issues in clinical ICT/PCT trials.

Patient's decision

In order to satisfy cancer patients and improve therapeutic outcomes, decision aids systems might also helpful in clinical therapeutics [27,28]. By providing decision aids service, cancer patients may be more cooperative and optimize for further clinical cancer therapeutics. This emerging technique may reach unexpected result in cancer treatments. This type of medical service might be added to future PCT/ICT items.

Artificial intelligence in ICT/PCT

Artificial intelligence techniques might sooner or later go into hospitals. This important cancer treatment trend must be noticed early. Since repetitive therapeutic work or variability of doctor's prescriptions will be superseded by computers, robots and other artificial intelligence technology. Don't miss this milestone work in future.

Economic consideration of ICT/PCT and cost-effective in clinical trials

Updating clinical cancer treatments is a difficulty and cost thing nowadays. Though it is a better way to benefit tumor growth/metastasis treatments by ICT/PCT, we still do not find any closer association between therapeutic benefits and therapeutic expenditures. A good balance between drug activity, toxicities and cost is the state-of-the-art system and new law of anticancer drug combination study. In order to do that, mathematical/computational network is the cheapest ways for simulations and decision-making. As a result, further work is needed in this avenue.

Since many ICT/PCT strategies are complementary with each other, two or three types of ICT strategies are seldom applicable in one cancer patient for economic considerations and technical feasibility. According to cancer patient' pathological situation and financial condition, we can optimize cancer therapeutics for patients in future [29-31]. However, selections of ICT/PCT must be based on cost-effective evaluations. Cost-effective study of drug combination and biotherapies, such as gene therapy or antibody therapy is main parts of ICT/PCT owing to their relatively higher therapeutic costs. Considering more than \$10,000 expenditure of common cycles of drug combination/biotherapy, the cost of new types of ICT/PCT strategic options may be much less if we can maintain the diagnostic fee to \$30 - 5,000. ICT/PCT of less than 5,000 USD is certainly cost-effectiveness. After new cost-effective diagnostics, it will certainly increase the quality adjusted life year (QALY) of cancer patients, especially in some early stage of cancer or young cancer patients [29-31]. Almost each of presently used ICT/PCT strategies is cost-effective from broader definitions and possibly save the life of more cancer patients.

Comparisons of different items of ICT/PCT strategies

Since cancer treatments are far from our expectations and requirements-greatly elongation of cancer patient's survivals, ICT/PCT perfections is inevitable.

In order to improve currently available ICT/PCT strategies, new round of experimental/clinical campaign and ICT/PCT applications will be undergone. Presently, anticancer drug development is more suitable for human leukemia treatment and less effective to solid human tumors, especially to late-staged cancer patients. In future, more complex forms of ICT/PCT will be developed for cancer patients with solid tumors.

Among different types of ICT/PCT, which type of ICT/PCT strategy is the best? Each of them has its own advantages and disadvantage. At present no one type of ICT/PCT strategies is obviously advantageous over the others. In addition, no available ICT/PCT strategy has been well enough to significantly increase the patient's survivals comparing with conventional therapy. So, we desperately need some dramatic moves to create new generations of more matured ones by integrating the advantages of most ICT/PCT types. Although much effort has been made, main obstacles remain to be overcome. The most important drawbacks of these ICT/PCT strategies is there is almost no survival benefits in patients with noticeable metastatic nodules in spite of DST, PG and other item utilities [32-35]. But it can be a future miracle if we can perfect them into a successful one. So, are we ready for that yet? [81].

Disease information modality

Cancer is a disease of great diversity of tumor genetic alterations or molecular abnormalities. Normally cancer can be categorized into 6 distinct hallmarks of pathologic processes (Table 1). The different ICT/PCT modalities are utilized for different hallmarks. It is proposed that different cancer hallmarks are suitable from different diagnostic modalities-PCT/ICT strategy.

Hallmarks of cancer	Suitable PCT/ICT items or strategy	
Sustaining proliferative signaling	DST, PG	
Resisting cell death	Biomarkers	
Inducing angiogenesis	Hormonal/assistant therapy, bioinformatics	
Evading growth suppressors	PG	
Enabling replicative immortality	PG, bioinformatics	
Invasion and metastasis	Drug combination, assistant therapy	

Table 1: Schematic outlook of associations between biology and pathology mechanisms and ICP/PCT strategies.

We have personally categorized these relationships into table 1. Following sectors will discuss other important items of PCT/ICT strategies.

Table 2 and 3 represents the technical cores of different types of ICT/PCT strategies nowadays (Table 2 and 3). From our vision, an integrated ICT/PCT strategy/paradigm will be designed from essence of these core technologies. Future biomedical efforts will offer these details and invite new items of ICT/PCT strategies with integrated technique and cutting-edge medical knowledge.

Major strategy	DNA	RNA	Macromolecules	Bioassay
DST	$\sqrt{}$	$\sqrt{}$	$\sqrt{}$	+++
PG	+++	+	+	√
Bioinformatics	+	++	+++	√
Metastasis	$\sqrt{}$	$\sqrt{}$	++	++
Drug combinations	√	√	+	++
Hormonal/assistant	√	√	++	++

Table 2: The associations between PCT strategies and biological suitability.

Symbolic Meaning + Some Suitability; ++ Good Suitability; +++Excellent Suitability; $\sqrt{\text{Needs Future Improvements.}}$

ICT strategies	Core techniques	
DST	Cell number or viability	
PG	SNP	
Metastasis condition	Tomography	
Bioinformatics	High-throughput techniques (omics)	
Drug combinations	Pharmacology	
Hormonal/assistant therapy	Drug characters	

Table 3: Technical cores of different ICT/PCT strategies nowadays.

Next Generation of ICT/PCT Strategies

Integration is the key

One decade ago, we proposed that integrated ICT/PCT systems would be a future trend [5]. After more than one decade, we reiterate our past argument in this article with newest twist.

The good ICT/PCT strategies must contain multiple functionality/therapeutic information-including cancer property revealing, drug types/responsibility, mathematical/computational information treatments [82-86] and decision-aids system [27,28] to individual patients. Thus, future strategy of ICT/PCT must at least contain a process of drug response (pharmacology), tumor pathogenesis information (oncology) and others [87,88]. In the past, pharmacology (DST and drug combination) and tumor characters and pathologic processes (tumor bioinformatics, biomarkers and PG) are separately determined in the clinic. This situation needs to be improved. Certainly, more dramatic technical innovations, such as next generation sequence (NGS) [89] must be incorporated into new generations of ICT/PCT systems and strategies in future. Among these trends, computational network, patient's decision aids and artificial intelligence must be updated and integrated into one system of ICT/PCT.

Afterwards, ICT/PCT needs to be less and less moneys and multi-levels of medical/pharmacological information. New techniques might change the landscape or blueprint of ICT/PCT study and application scenarios. Some longstanding questions of cancer biology and pathology, such as relationship between cancer heterogeneity and different drug combination therapy might provide new foundation. High-resolution and lower cost ICT/PCT from technical innovations and medical/pharmacologic advancements might create growing usefulness of ICT/PCT strategies in the future (Figure 3).

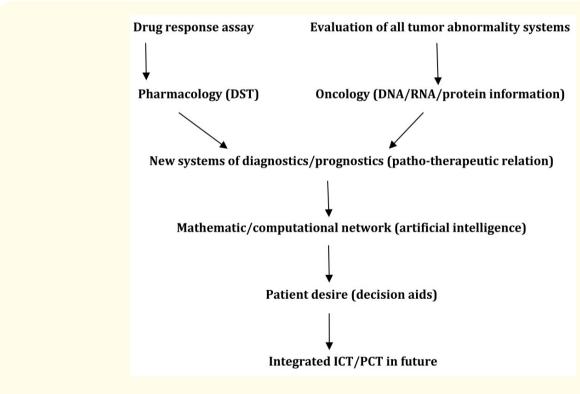


Figure 3: General scheme of ICT/PCT strategy developments in future.

Anticancer drug combination updating

Drug combination optimum strategy is too complicated to be easily mastered by clinical doctors. Though drug combination is a common way to enhance patients' therapeutic outcomes and survivals, there is still much room for updating and improving in both experimental and clinical investigations. In the past, clinical anticancer drug combination is based on doctors' judgment without in-depth therapeutic mechanism supporting. Clinical cancer combination therapies are generally empirical, statistics-oriented or past references rather than well-defined diagnostic/pharmacological association study. Finding new laws regarding anticancer drug combination (efficacy vs toxicity) must be established in future-central dogma establishments [74]. Due to highly complication from medical/pharmacologic situations, we cannot as usual overlook it [75].

The establishment of central dogma for drug combination is no easy task. In order to find this information, every possibility of drug combination must be undergone. Nowadays, approximately 178 anticancer drugs have been licensed worldwide [74-75],

According to mathematic equation (calculation for 2 - 3 anticancer drug combinations); (equation 1 - 2)

```
C2 = (178 \times 177)/(1 \times 2) = 15,753 (equation 1)
C3 = (178 \times 177 \times 176)/(1 \times 2 \times 3) = 924,176 (equation 2)
```

At the beginning of this study, we are awed with high numbers of drug combination possibility (Equation 1 and 2). However, when we think about the number of cancer morbidity worldwide, we are relieved suddenly [78]. Our conclusion is that only global coordination of this study can facilitate and master the core of drug combination practice (central dogma) sooner or later.

Conclusion

ICT/PCT is a difficult approach [90]. In order to in depth understand cancer pathology and therapy for individual patients, well-defined, prospective, retrospective and double-blind ICT/PCT clinical studies and clinical applications are urgently needed. We look forward to some integrated ICT/PCT strategies to be established from empirical to scientific-guided systems. From our perspective of ICT/PCT innovations, integration is the key!

Acknowledgements

This work was funded by Shanghai Science and Technology Foundation of High Educations 97A49.

Competing Interesting

Authors have declared that no competing interests exist.

Bibliography

- 1. Siegel RL., et al. "Cancer stastistics 2017". CA- Cancer Journal for Clinicians 67.1 (2017): 7-30.
- 2. Ali I., et al. "Social aspects of cancer genesis". Cancer Therapy 8.6 (2011): 6-14.
- 3. Nowell PC. "The clonal evolution of tumor cell populations". Science 194.4260 (1976): 23-28.
- 4. Hanahan D and Weinberg RA. "Hallmarks of cancer, the next generation". Cell 144.5 (2011): 646-674.
- 5. Lu DY., *et al.* "Individualized cancer chemotherapy integrating drug sensitivity tests, pathological profile analysis and computational coordination-an effective strategy to improve clinical treatment". *Medical Hypotheses* 66.1 (2006): 45-51.
- 6. Lu DY., et al. "Individualized cancer chemotherapy". Hypotheses in Clinical Medicine Ed, Shoja MM, Agutter PS, Tubbs RS, Ghanei M, Ghabili K, Harris A, Loukas M. chapter 13, Nova Science Publisher. US (2012): 199-216.

- 7. Lu DY., et al. "Personalized cancer therapy, a perspective". Clinical Experimental Pharmacology 4.2 (2014) 153.
- 8. Lu DY, "Personalized cancer therapy, a perspective". *International Journal of Pharmacy Practice and Drug Research* 4.2 (2014): 108-118.
- 9. DY. "Personalized cancer chemotherapy, an effective way for enhancing outcomes in clinics". Woodhead Publishing, Elsevier, UK (2014).
- 10. Lu DY, et al. "Individualized cancer therapy". Innovations in Pharmaceuticals and Pharmacotherapy 2.4 (2014): 458-469.
- 11. Lu DY, et al. "Anticancer drug sensitivity testing, a historical review and future perspectives". Current Drug Therapy 10.1 (2015): 44-55.
- 12. Lu DY. "Drug sensitivity testing. Personalized Cancer Chemotherapy, An Effective Way for Enhancing Outcomes in Clinics". Ed, Lu DY, Chapter 2, Woodhead Publishing, Elsevier, UK (2014): 5-12.
- 13. Volm M and Efferth T. "Prediction of cancer drug resistance and implications for personalized medicine". Frontiers in Oncology 5 (2015): 282.
- 14. Lu DY, et al. "Cancer bioinformatics, its impacts on cancer therapy". Metabolomics 5.2 (2015): e133.
- 15. Ocana A and Pandiella A. "Personalized therapies in the cancer "omics" era". Molecular Cancer 9 (2010): 202.
- Stransky B and Galante P. "Application of bioinformatics in cancer research". An IMICS Perspective on Cancer Research (2010): 211-233.
- 17. Lu DY. "Individualized cancer chemotherapy via cancer biomarkers or bioinformatics detecting". Personalized Cancer Chemotherapy, An Effective Way for Enhancing Outcomes in Clinics. Ed, Lu DY, Chapter 3, Woodhead Publishing, Elsevier, UK (2014): 13-20.
- 18. Lu DY., et al. "Cancer bioinformatics for update anticancer drug developments and personalized therapeutics". Reviews on Recent Clinical Trials 12.2 (2017): 101-110.
- 19. Meyer UA. "Pharmacogenetics-five decades of therapeutic lessons from genetic diversity". *International Journal of Pharmacy Practice and Drug Research* 5.9 (2004): 669-676.
- 20. Lu DY. "Pharmacogenetics. Personalized Cancer Chemotherapy, An Effective Way for Enhancing Outcomes in Clinics. Ed, Lu DY, Chapter 4, Woodhead Publishing, Elsevier, UK (2014): 21-28.
- 21. Lu DY, et al. "Pharmacogenetics of cancer therapy: breakthroughs from beyond? Future Science OA 1.4 (2015): 15.
- 22. Lu DY, et al. "New insights into individualized antimetastatic therapy". Advanced Techniques in Biology and Medicine 1.1 (2013): 106.
- 23. Lu DY. "Individualized antimetastatic therapy. Personalized Cancer Chemotherapy, An Effective Way for Enhancing Outcomes in Clinics". Ed, Lu DY, Chapter 5, Woodhead Publishing, Elsevier, UK (2014): 29-36.
- 24. Lu DY. "Drug combinations. Personalized Cancer Chemotherapy, An Effective Way for Enhancing Outcomes in Clinics. Ed, Lu DY, Chapter 6, Woodhead Publishing, Elsevier, UK (2014): 37-42.
- 25. Lu DY. "Assistant chemotherapy. Personalized Cancer Chemotherapy, An Effective Way for Enhancing Outcomes in Clinics. Ed, Lu DY, Chapter 7, Woodhead Publishing, Elsevier, UK (2014): 43-48.

- 26. Lu DY., et al. "Old theories revisited on cancer assistant therapy". International Journal of Medical Research and Health Sciences 1 (2014): 50-57.
- 27. O'Connor A. "Using decision aids to help patients navigate the "grey zone" of medical decision-making". CMAJ 176.11 (2007): 1597-1598.
- 28. Holbrook A., et al. "Influence of decision aids on patient preferences for anticoagulant therapy: a randomized trial". CMAJ 176.11 (2007): 1583-1587.
- 29. Lu DY., et al. "Cost-effectiveness considerations of individualized cancer chemotherapy". Advances in Pharmacoepidemiology and Drug Safety 2.5 (2013): e121.
- 30. Naeim A and Keeler EB. "Is adjuvant therapy for older patients with node (-) early breast cancer cost-effective?". *Critical Review in Oncology/Hematology* 53.1 (2005): 81-89.
- 31. Lu DY. "Cost-effectiveness consideration. Personalized Cancer Chemotherapy, An Effective Way for Enhancing Outcomes in Clinics". Ed, Lu DY, Chapter 8, Woodhead Publishing, Elsevier, UK (2014): 49-59.
- 32. Talmadge JE and Fidler IJ. "The biology of cancer metastasis: historical perspective". Cancer Research 70.14 (2010): 5649-5669.
- 33. Valastyan S and Weinberg RA. "Tumor metastasis: molecular insights and evolving paradigms". Cell 147.2 (2011): 275-292.
- 34. Lu DY, et al. "Cancer metastases and clinical therapies". Cell and Developmental Biology 1.4 (2012): e110.
- 35. Lu DY., et al. "Cancer Metastasis treatments". Current Drug Therapy 8.1 (2013): 24-29.
- 36. Lu DY., et al. "Antimetastatic activities and mechanisms of bisdioxopiperazine compounds". Anti-Cancer Agent Medicinal Chemistry 10.7 (2010): 564-570.
- 37. Mina LA and Sledge GW. "Rethinking the metastatic cascade as a therapeutic target". *Nature Reviews Clinical Oncology* 8.6 (2011): 325-332.
- 38. Paez-Ribes M., et al. "Antiangiogenic therapy elicits malignant progression of tumors to increased local invasion and distant metastasis". Cancer Cell 15.3 (2009): 220-231.
- 39. Ebos JML, *et al.* "Accelerated metastasis after short-term treatment with a potent inhibitor of tumor angiogenesis". *Cancer Cell* 15.3 (2009): 232-239.
- 40. Lu DY., et al. "Antimetastatic activities and mechanisms of action among Bisdioxopiperazine compounds. Pharmaceutical Formulation and Medicinal Chemistry: Mechanisms, Developments and Treatments". Ed. Bruce Moore. chapter 2, Nova Science Publishing, US (2016): 73-106.
- 41. Lu DY, et al. "Anticancer drug development, system updating and global participations". Current Drug Therapy 12.1 (2017): 37-45.
- 42. Lu DY, et al. "Anticancer drug development, getting out from bottleneck". Journal of Medicinal Chemistry 7.1 (2017): 423.
- 43. Gupta SC., et al. "Cancer drug discovery by repurposing: teaching new tricks to old dogs". *Trends in Pharmacological Sciences* 34.9 (2013): 507-517.
- 44. Lu D., et al. "Structural aberration of cellular sialic acids and their functions in cancer". Journal of Shanghai University (English Edition) 5.2 (2001): 164-170.

- 45. Varki NM and Varki A. "Diversity in cell surface sialic acid presentations: implications for biology and disease". *Lab Investigation* 87.9 (2007): 851-857.
- 46. Varki A. "Sialic acids in human health and disease". Trends in Molecular Medicine 14.8 (2008): 351-360.
- 47. Lu DY, et al. "Antimetastatic therapy targeting aberrant sialylation profiles in cancer cells". Drug Therapy Studies 1.1 (2011): e12.
- 48. Lu, D.Y., et al. "Development of antimetastatic drugs by targeting tumor sialic acids". Scientia Pharmaceutica 80.3 (2012): 497-508.
- 49. Bull C., *et al.* "Targeting aberrant sialylation in cancer cells using a fluorinated sialic acid analog impairs adhesion, migration, and in vivo tumor growth". *Molecular Cancer Therapeutics* 12.10 (2013): 1935-1946.
- 50. Xi YC and Lu DY. "The relationship between tumor sialic acids and neoplasm metastases". Cell and Developmental Biology". 1.3 (2012): e106.
- 51. Lu DY., *et al.* "Inhibitions of some antineoplastic drugs on serum sialic acid levels in mice bearing tumors". *Scientia Pharmaceutica* 81.1 (2013): 223-231.
- 52. Lu DY, *et al.* "Different spontaneous pulmonary metastasis inhibitions against Lewis lung carcinoma in mice by bisdioxopiperazine compounds of different treatment schedules". *Scientia Pharmaceutica* 78.1 (2010): 13-20.
- 53. Marx V. "Tracking metastasis and tracking cancer". Nature 494 (2013): 131-136.
- 54. Reuben JM., *et al.* "Circulating tumor cells, disease progression, and survival in metastatic breast cancer". *The New England Journal of Medicine* 351 (2004): 781-791.
- 55. Lu DY, et al. "Effects of cancer chemotherapy on the blood fibrinogen concentrations of cancer patients". *Journal of International Medical Research* 28.6 (2000): 313-317.
- 56. Lu DY., et al. "Effects of anticancer drugs on the binding of 125I-fibrinogen to two leukemia cell lines in vitro". Journal of International Medical Research 32.5 (2004): 488-491.
- 57. Lu DY., *et al.* "Treatment of solid tumors and metastases by fibrinogen-targeted anticancer drug therapy". *Medicinal Hypotheses* 68.1 (2007): 188-193.
- 58. Lu DY., et al. "Relationship between blood fibrinogen concentration and pathological features of cancer patients: a 139-case clinical study". OnLine Journal of Biological Sciences 7.1 (2007): 8-11.
- 59. Bobek V. "Anticoagulant and fibrinolytic drugs-possible agents in treatment of lung cancer?". *Anticancer Agents in Medicinal Chemistry* 12.6 (2012): 580-588.
- 60. Lu DY., et al. "Tumor fibrin/fibrinogen matrix as a unique therapeutic target for pulmonary cancer growth and metastases". Pulmonology and Clinical Research 3.1 (2015): 1027.
- 61. Rothwell P., et al. "Effect of daily aspirin on long-term risk of death due to cancer: analysis of individual patient data from randomized trials". *Lancet* 377.9759 (2011): 31-41.
- 62. Thiery JP, et al. "Epithelial-mesenchymal transitions in development and disease". Cell 139.5 (2009): 871-890.
- 63. Kalluri R and Weinberg RA. "The basics of epithelial-mesenchymal transition". *The Journal of Clinical Investigation* 119.6 (2009): 1420-1428.

- 64. Tsai JH., et al. "Spatiotemporal regulation of epithelial mesenchymal transition is essential for squamous cell carcinoma metastasis". Cancer Cell 22.6 (2012): 725-736.
- 65. Ocana OH., et al. "Metastatic colonization requires the repression of the epithelial-mesenchymal transition inducer Prrx1". Cancer Cell 22.6 (2012): 709-724.
- 66. Brabletz T. "EMT and MET in metastasis: where are the cancer stem cells?". Cancer Cell 22.6 (2012): 699-701.
- 67. Van Denderen BJW and Thompson EW. "The to and fro of tumour spread". Nature 493.7433 (2013): 487-488.
- 68. Lu DY, et al. "Cancer metastasis, a clinical dilemma for therapeutics". Current Drug Therapy 11.2 (2016): 163-169.
- 69. Pavelic SK., et al. "Metastasis: new perspectives on an old problem". Molecular Cancer 10 (2011): 22.
- 70. Olson BM and McNeel DG. "Monitoring regulatory immune responses in tumor immunotherapy clinical trials". *Frontier in Oncology* 3 (2013): 109.
- 71. Lu DY., et al. "Combination chemical agents with biological means in cancer therapy". Research and Reviews in BioScience 7.4 (2013): 153-155.
- 72. Tipping AJ and Melo JV. "Imatinib mesylate in combination with other chemotherapeutic drugs: In vitro studies". *Semin Hematol* 40.2 (2003): 83-91.
- 73. Druker B. "Imatinib alone and in combination for chronic myeloid leukemia". Seminars in Hematology 40.1 (2003): 50-58.
- 74. Lu DY, et al. "Drug combinations in cancer treatment". Clinical Experimental Pharmacology 3 (2013): 134.
- 75. Lu DY, et al. "Anticancer drug combination, how far we can go through?". Anti-Cancer Agents in Medicinal Chemistry 17.1 (2017): 21-28.
- 76. Lu DY., et al. "Anticancer drug combinations, a big momentum is needed". Metabolomics 5.3 (2015): e139.
- 77. Lu DY., et al. "Anticancer drug combinations: the next medical challenge". *Innovations in Pharmaceuticals and Pharmacotherapy* 3.3 (2015): 637-649.
- 78. Lu DY, et al. "Drug combination in clinical cancer treatments". Reviews on Recent Clinical Trials (2017).
- 79. Lu DY, et al. "Anticancer drug combinations, studies from different pathways". Cell and Developmental Biology 4.5 (2015): 166.
- 80. Lu DY., et al. "Anticancer drug combinations, studies for all possibilities". Advances in Pharmacoepidemiology and Drug Safety 5.1 (2016) e138.
- 81. Lu DY, et al. "Individualized cancer chemotherapy, are we ready for that yet?" Metabolomics 2 (2012): e113.
- 82. Komarova NL. "Mathematical modeling of tumorigenesis, mission possible". Current Opinion in Oncology 17.1 (2006): 39-43.
- 83. Khalil C. "System biology for cancer". Current Opinion in Oncology 17.1 (2006): 44-48.
- 84. Lu DY., et al. "Employing new mathematical models and equations to evaluate risk-benefit criteria of clinical therapeutics". Online Journal of Biological Science 7.1 (2007): 1-2.
- 85. Waterman MS. "Introduction to computational biology; maps, sequence and genomes". CRC Press, Taylor Francis Group LLC, US (2000).

- 86. Axelrod R and Hamilton WD. "The evolution of cooperation". Science 211 (1981): 1390-1396.
- 87. Lu DY, et al. "General topics in the field of personalized cancer therapy". Metabolomics 7.S1 (2017): e001.
- 88. Lu DY, et al. "Clinical cancer therapy, personalized chemotherapies". Journal of Cell and Developmental Biology 1.1 (2017): 5.
- 89. Lander ES. "Initial impact of the sequencing of the human genome". Nature 470 (2011): 187-197.
- 90. Span PN. "From eels to the importance of cancer biobanks". Future Science OA 1.4 (2015).

Volume 2 Issue 6 June 2018 ©All rights reserved by Da-Yong Lu., *et al.*