

A Comparative Study on Different Plasmid Isolation Procedures

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ABSTRACT

Plasmid DNA has several applications in the field of modern biotechnology. Despite the presence of several methods to remove contaminants and/ or to purify plasmid DNA from primary isolate, the first technique of plasmid isolation determines the quality of the final product. In this study, some popular methods of plasmid DNA isolation have been compared in terms of their rapidity, plasmid quality and ease of experiments. Three different STET buffers used in boiling lysis process were also compared to identify the best performing one. Highest quantity of RNA contaminants, as well as plasmid, was isolated in the alkaline extraction process. Best supercoiled plasmid percentage was isolated by boiling lysis STET buffer 2. The RE digestion was non-reactive/ improper in boiling lysis methods whereas it was perfect in alkaline lysis despite huge quantity of RNA contaminants. Alkaline lysis was proved to be the best one among all and with some further purification processes indicated, other processes can also be used efficiently.

Key words: Alkaline lysis, Boiling lysis, STET buffer, Plasmid isolation, Plasmid quality

Abbreviations-

BL1: Plasmid isolation by boiling lysis process with STET 1 buffer

BL2: Plasmid isolation by boiling lysis process with STET 2 buffer

BL3: Plasmid isolation by boiling lysis process with STET 3 buffer

INTRODUCTION

Plasmid is extra chromosomal DNA that exists mainly in prokaryotes like bacteria and archaea. However, extra chromosomal plasmids also exist in certain eukaryotic fungal species³⁰. These plasmids exhibit many interesting characteristics like antimicrobial resistance or fertility factor in the life of the host organism. Replication of plasmid does not depend upon the host chromosomal DNA

machinery but rather follows different approaches²⁶. Being a small (2-20 kb in average) independent DNA molecule, plasmid DNA has been the centre of the biotechnological studies for a long time. Plasmid DNAs are easily modifiable using genetic engineering approach and these can be maintained and replicated in a host bacterial organism (like *Escherichia coli*).

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In modern days, plasmid DNAs are used in thousands of ways including sequencing, gene expression, promoter analysis, gene therapy, DNA vaccine, gene knockdown/ knockout, CRISPR-Cas, production of transgenic organism and many more. Successes of these applications depend upon the quality of the plasmid DNA. Plasmids exist in a complex topologically intertwined supercoiled structure inside the cell. Due to the mechanical or shear stress applied during extraction, the plasmid comes in some other forms like open circular, linear, covalently bounded (figure 1). But, the supercoiled plasmids are mostly accepted by cells during transformation or transfection processes. The final plasmid should be free from any traces of contaminants like degraded RNA, genomic DNA, bacterial proteins and endotoxins. There are several approaches to remove these contaminants like use of chloride salts to remove high molecular weight RNA^{7,29}, selective precipitation of supercoiled plasmid^{21,25,29,32}, filtration¹² and chromatographic techniques¹¹. Though all these techniques are standardized over time to work efficiently, the primary technique of

plasmid isolation determines the quality of the final product.

In this study, we have compared some popular methods of plasmid extraction in terms of their rapidity, plasmid quality and ease of experiments. Alkaline extraction is the most popular method of plasmid extraction till date. In this process, the bacterial cell is ruptured by exposing it to a highly alkaline solution along with detergent. The gDNA and cellular proteins are selectively precipitated by addition of an acidic neutralization solution while the topologically intertwined plasmids are retained in the aqueous phase^{2,26}. Boiling lysis is another popular method of plasmid isolation in which cells are ruptured with a combination of detergent, lysozyme and a heat shock. The chromosomal DNA remains attached to bacterial proteins and is precipitated by centrifugation while the plasmid remains in the supernatant¹⁸. The lysis buffer used in boiling lysis method somewhat differs in available different literature. In this study, we have also compared all the different compositions to find out the best working one.

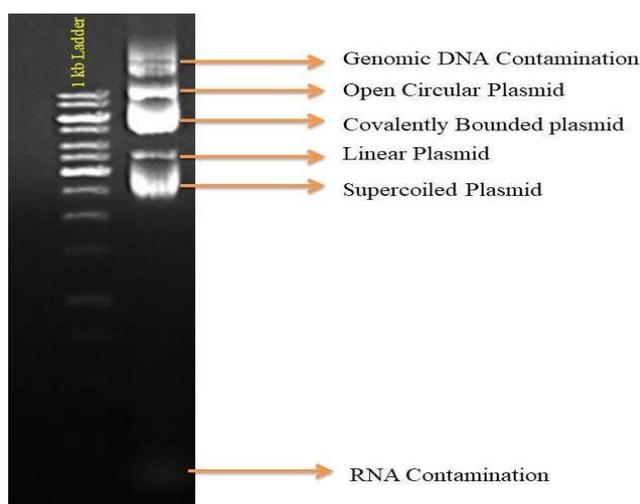


Figure 1: Different contaminants and isoforms of plasmids after plasmid extraction process. Photograph obtained after 1% agarose gel electrophoresis with voltage of 5 volt/cm of gel for 1.5 hours.

MATERIAL AND METHODS

Bacterial culture

E.coli (strain DH5 α), harbouring a kanamycin resistance gene containing plasmid was cultured in Luria bertani broth with 50 μ g/ ml of kanamycin selection pressure at 37°C in a shaking incubator at 160 rpm overnight (16 hours). The final OD₆₀₀ of the culture was 1.12. The same wet weight of bacterial pellet (20 mg) was taken in all the experiments.

Plasmid isolation

The plasmid was extracted mainly by two methods. The alkaline extraction procedure was carried out according to Sambrook *et al.*²⁶. The boiling lysis method was carried out using three different lysis buffers (named STET buffer 1, 2 & 3). The composition of the buffer system is given in table no. 1. As a positive control, plasmid was also isolated using Qiagen miniprep kit following the manufacturer's protocol. All the experiments were carried out in triplicates.

Table 1: Comparison of different STET buffer compositions

Buffer	Composition	Reference
STET 1	8% Sucrose, 5% Triton X-100, 50 mM EDTA (pH-8), 50 mM Tris-Cl (pH-8). The solution is autoclaved and stored at 4°C.	18
STET 2	8% Sucrose, 0.5% Triton X-100, 50mM EDTA (pH-8), 10mM Tris-Cl (pH-8). The solution is autoclaved and stored at 4°C.	This study
STET 3	10mM Tris-Cl (pH-8), 0.1M NaCl, 1mM EDTA (pH-8), 5% v/v Triton X-100. pH of STET is 8.0 after all the ingredients are added. No need of sterilize before use.	26

Procedure for plasmid isolation in boiling lysis was carried out according to Sambrook *et al.*²⁶ with some minor modifications. Briefly, 20 mg (wet weight) of bacterial pellet was washed with PBS and resuspended in 350 μ l of STET buffer by slow vortexing. Resuspended solutions were incubated in ice for 5 minutes. 25 μ l of freshly prepared lysozyme solution (10 mg/ ml in 10mM Tris-cl, pH 8.0) was added to the resuspended bacterial solution. After thorough mixing, the tubes were kept in a boiling water bath (100°C) for exactly 40 seconds and immediately kept on ice for 2 minutes. The tubes were then centrifuged at 14000 rpm for 20 minutes at 4°C. The resultant supernatant was decanted in another tube, rather than pipetting out because pipetting can sometimes result in entry of viscous protein containing liquid. But, decanting is only recommended after application of centrifugation time and speed mentioned above. 0.6 volume of room temperature isopropanol was added to the supernatant, incubated at room temperature for 15 minutes and centrifuged at 12000 rpm for 10 minutes at room temperature. The

supernatant was discarded and the pellet was washed twice with cold 70% alcohol. Centrifugation after alcohol wash was avoided. Lastly, the pellets were resuspended in 30 μ l of TE buffer.

Plasmid quality determination

The plasmid quality was determined depending upon their plasmid yield, 260/280 ratio, RNA contamination, genomic DNA contamination, supercoiled plasmid percentage and RE digestion. Plasmid yield and 260/280 ratio was determined by spectrophotometric method in nan drop. RNA contamination was estimated by 1% agarose gel electrophoresis. Genomic DNA contamination percentage and supercoiled plasmid DNA percentage was determined using my Image Analysis software (Thermo Scientific, USA) by analysing the gel image. Double RE digestion [using Xho1 and Sph1 (Thermo Scientific, USA)] was performed to identify the presence of enzyme inhibitory contaminants in the isolate. RE digestion was performed at 37°C for 3 hours without RNase treatment.

RESULTS

Plasmid yield

The boiling lysis procedure took almost 1 hour 30 minutes whereas the alkaline lysis method took nearly 2 hours and the Qiagen miniprep kit took 30 minutes to be completed. In BL1, BL2, BL3, alkaline lysis and Qiagen miniprep kit, the total plasmids extracted are 33.6 μ g,

62.21 μ g, 57.7 μ g, 86 μ g and 7.6 μ g respectively. Among the traditional methods, the best 260/280 ratio (near to 1.8) was obtained in BL1 method. The supercoiled percentage was highest in BL2 method (34.32%). Alkaline lysis and BL3 methods showed the highest quantity of genomic DNA contamination (shown in figure 2). The other quality data are noted in table 2.

Table 2: Quality comparison of plasmid isolated by different processes

Processes	Boiling Lysis	Boiling Lysis	Boiling Lysis	Alkaline	Qiagen
Quality	STET 1	STET 2	STET 3	Lysis	Miniprep
Plasmid yield (μ g)	33.6	62.1	57.7	86	7.6
260/280	1.935	2.075	2.04	2.168	1.801
Supercoiled %	33.4	34.32	32.66	27	30
Genomic DNA contamination	Low	Low	High	High	Low
RE digestion	Negative	Negative	Partial	Positive	Positive

RE digestion

The RE digestion was performed using Xho1 and Sph1 enzymes at 37°C for 3 hours. The expected fragment sizes were 1.2 kb and 2.0 kb. No bands were seen in case of BL1 and BL2 plasmids after RE digestion. In the case

of BL 3 plasmid, an improper RE was observed which provided faint bands of expected sizes. The plasmids isolated by alkaline lysis and Qiagen miniprep kit were digested perfectly (figure 3).

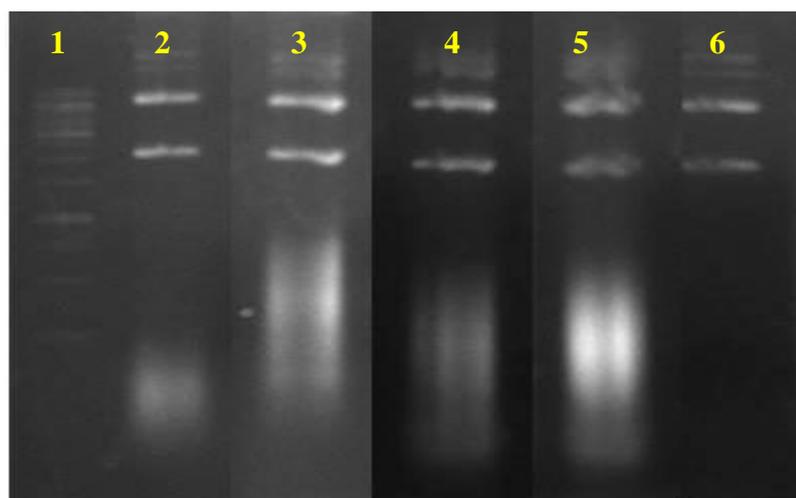


Figure 2: Agarose gel electrophoresis image of plasmids isolated by different procedures. 1 kb DNA ladder (lane 1); plasmid isolated by BL1 method (lane 2); plasmid isolated by BL2 method (lane 3); plasmid isolated by BL3 method (lane 4); plasmid isolated by alkaline lysis procedure (lane 5); plasmid isolated by Qiagen miniprep kit (lane 6).

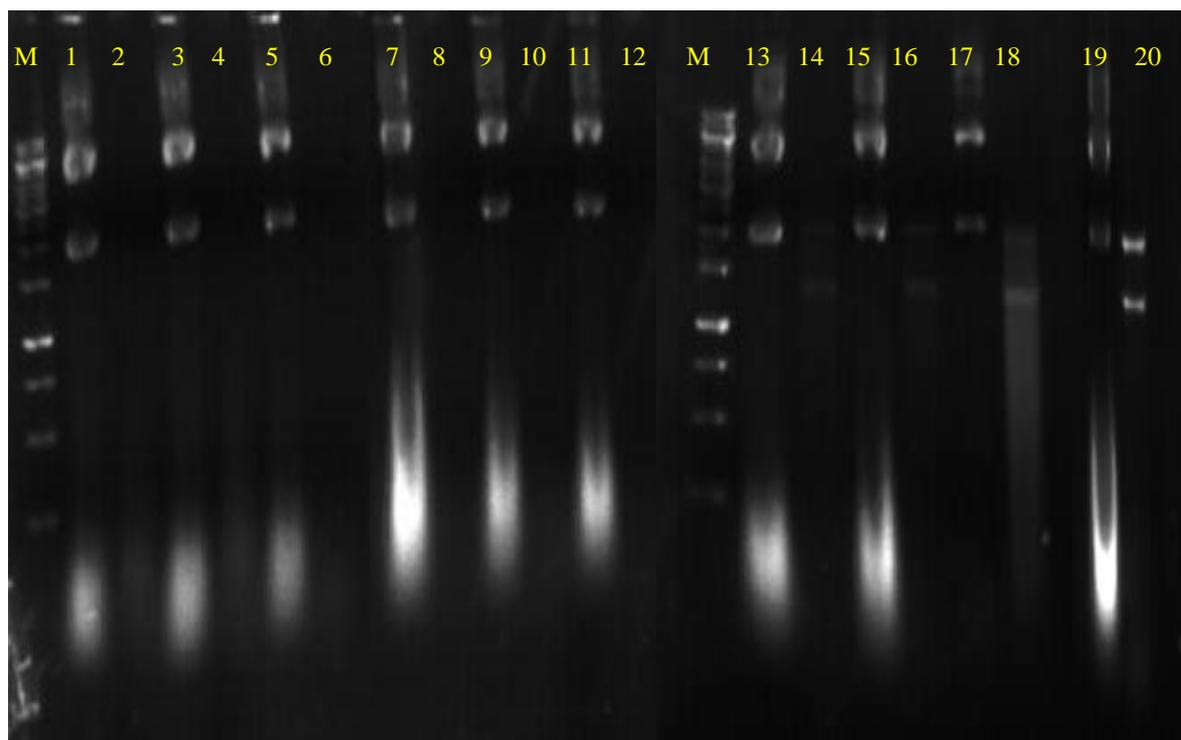


Figure 3: RE digestion of plasmids.

1 kb DNA ladder (lane M); plasmids isolated by BL1 (lane 1, 3, 5); RE of plasmids isolated by BL1 (lane 2, 4, 6); plasmids isolated by BL2 (lane 7, 9, 11); RE of plasmids isolated by BL2 (lane 8, 10, 12); plasmids isolated by BL3 (lane 13, 15, 17); RE of plasmids isolated by BL3 (lane 14, 16, 18); plasmid isolated by alkaline lysis (lane 19); RE of plasmid extracted by alkaline lysis (lane 20).

DISCUSSIONS

In the current study, we compared some popular methods for plasmid isolation to find out the most efficient one in terms of rapidity, plasmid quality and ease of experiment. Although the Qiagen miniprep kit follows the basic principle of alkaline lysis method, it is far more rapid than the traditional one because it contains RNase in the resuspension buffer solution and it omits most of the incubation steps to be followed in the traditional process. The boiling lysis method is sufficiently faster (1.5 hours) than the alkaline lysis method (2 hours). However, the former one needs instruments like boiling water bath which may not be a common instrument in common molecular biology laboratory. Furthermore, the time required for alkaline lysis process can be reduced by minimizing or omitting some of the incubation steps.

The spectrophotometric highest yield of plasmid was obtained in alkaline lysis

method which was confirmed by agarose gel electrophoresis image (figure 2). Subsequently, the lowest yield was in Qiagen miniprep kit as the capacity of the silica column used in the kit is limited to 10 μ g only. BL 1 yielded somewhat lower yield than the other two methods in spectrophotometric assay, but the result was not in line with the gel image (figure 2). The purity (260/280 ratio) of the isolated plasmid can solve this biases as the purity of the BL1 plasmid was somewhat higher than the others (table 2). It is possible that the contaminants in BL2 and BL3 plasmids probably contributed in misreading the actual absorption. The alkaline lysis isolated plasmids were the most RNA contaminated ones which can be seen by its high 260/280 ratio (2.168) and the large smear in gel image (figure 2). The boiling lysis method is not recommended when the copy number of plasmid is low as its efficiency is lower than alkaline lysis^{9,26}. Our result also

indicated lower yield in boiling lysis procedures. The yield was sufficiently high because the plasmid used by us has high copy number pUC origin of replication.

The supercoiled plasmid percentage was higher in all the boiling lysis methods with BL2 being the highest (34.32%). In contrast, the alkaline lysis and the kit method both possessed a lower percentage of supercoiled plasmid (table 2). The cell lysis period in alkaline lysis period and mixing pattern after addition of lysis buffer are very crucial steps for good quality supercoiled plasmid isolation²⁶. The standard incubation period for lysis in alkaline lysis is 5 minutes. However, the lysis time can vary depending upon the strain of bacteria, amount of bacteria and mixing. Careful standardization of lysis time according to the mentioned factors can help to yield more supercoiled DNA.

BL1 and BL2 plasmids failed to be digested properly in restriction analysis whereas the BL3 plasmids digested partially. The bacterial species that release carbohydrates in presence of detergent, lysozyme and heat (HB101) and the endonuclease A (*endA+*) strains (HB101, JM100) are not suitable for plasmid extraction by boiling lysis procedure^{9,26}. Contamination by endonuclease is not completely removed during boiling and can result in complete plasmid degradation when RE digestion is performed in presence of Mg²⁺ ion⁹. Although DH5 α is neither a carbohydrate releasing or *endA+* strain, it is suspected that the presence of other endonucleases has caused the complete disappearance of the plasmid. The BL3 method contained somewhat less quantity of endonuclease and showed improper RE digestion. In contrast, the plasmid isolated by alkaline lysis method was properly digested by restriction enzymes, indicating that although the isolate had a huge quantity of RNA contamination; it was completely free from endonucleases or any other enzyme inhibitory substances.

The problem of contamination can be avoided by certain additional steps in all the traditional processes. An additional phenol-

chloroform extraction step can be implemented after the first supernatant isolation in boiling lysis procedure to get rid of endonuclease problem⁹. The contamination of RNA in alkaline extraction can be completely removed by a brief treatment with RNase A. But, both the phenol and RNase A can be unsuitable for in vivo applications like gene therapy or DNA vaccination^{4,13} because the phenol is a corrosive agent and RNase is of bovine origin and can carry unnecessary harmful viruses.

There are several methods for selective purification of supercoiled plasmid DNA after isolation. Density gradient centrifugation using cesium chloride and ethidium bromide was one of the first methods of purifying supercoiled plasmid DNA¹⁴. However, the yield is low in this technique and both cesium chloride and ethidium bromide are toxic for cells. Because of these restrictions, density gradient centrifugation is rarely used for plasmid purification now. Removal of RNA contamination from plasmid DNA solution has been successfully carried out by selective precipitation with calcium chloride²⁹, lithium, ammonium acetate¹⁹ etc. or by filtration techniques like tangential flow filtration^{12,16,24,31}. Selective precipitation of supercoiled plasmid DNA can be achieved by use of polyethylene glycol (PEG)²⁹, cetyltrimethylammonium bromide (CTAB)²¹, polyamines²⁵, poly (*N,N*-dimethyldiallylammonium) chloride³² etc. Chromatographic techniques are the modern ones used to selectively purify supercoiled plasmid DNA. Although many types of chromatography including size exclusion^{6,22}, hydrophobic interaction⁸, hydroxyapatite¹⁵, thiophilic interaction²⁷, anion exchange¹¹ are used, anion exchange chromatography proved to be the most successful and efficient one. Techniques like monolith columns^{1,5,17,20} and chromatographic capsules¹⁰ are being increasingly popular over traditional chromatographic columns. Mostly, combinations of different techniques are applied for ease of purification. For small scale applications, homemade filtration column²⁸ or

homemade silica columns^{3,23} can be efficiently used.

From this study, we conclude that though the alkaline lysis process produces a lot of RNA contamination, the plasmid yield is way higher than other procedures tested. Moreover, the alkaline lysis plasmid is completely free from any enzyme inhibitory compounds. The boiling lysis procedure can be unsuitable for enzymatic reactions without any additional treatment. Any purification technique or combination of purification techniques mentioned above can be used effectively to further purify the DNA for use in other applications.

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Conflict of interest

The authors declare that they do not have any conflict of interest.

REFERENCES

1. Benčina, M., Podgornik, A. and Štrancar, A., Characterization of methacrylate monoliths for purification of DNA molecules. *Jour of separ sci*, **27(10-11)**: 801-810 doi: <https://doi.org/10.1002/jssc.200401784> (2004).
2. Bimboim, H. C. and Doly. J., A rapid alkaline extraction procedure for screening recombinant plasmid DNA. *Nucl acids res*, **7(6)**: 1513-1523 doi: <https://doi.org/10.1093/nar/7.6.1513> (1979).
3. Borodina, T. A., Lehrach, H. and Soldatov, A. V., DNA purification on homemade silica spin-columns. *Analyt biochem*, **321(1)**: 135-137 doi: [https://doi.org/10.1016/S0003-2697\(03\)00403-2](https://doi.org/10.1016/S0003-2697(03)00403-2) (2003).
4. Brand, E., Ralla, K. and Neubauer. P., Strategies for plasmid DNA production in Escherichia coli. *Biopharm Produc Tech*, **1**: 1-41 doi: <https://doi.org/10.1002/9783527653096.ch1> (2012).
5. Branovic, K., Forcic, D., Ivancic, J., Strancar, A., Barut, M., Gulija, T. K. and Mazuran. R., Application of short monolithic columns for fast purification of plasmid DNA. *Jour of Chromato B*, **801(2)**: 331-337 doi: <https://doi.org/10.1016/j.jchromb.2003.11.035> (2004).
6. Bywater, M., Bywater, R. and Hellman, L., A novel chromatographic procedure for purification of bacterial plasmids. *Analyt biochem*, **132(1)**: 219-224 doi: [https://doi.org/10.1016/0003-2697\(83\)90451-7](https://doi.org/10.1016/0003-2697(83)90451-7) (1983).
7. Chakrabarti, A., Sitaric, S. and Ohi, S., A procedure for large-scale plasmid isolation without using ultracentrifugation. *Biotech and appl biochem*, **16(2)**: 211-215 doi: <https://doi.org/10.1111/j.1470-8744.1992.tb00224.x> (1992).
8. Diogo, M. M., Queiroz, J. A., Monteiro, G. A. and Prazeres, D. M., Separation and analysis of plasmid denatured forms using hydrophobic interaction chromatography. *Analyt biochem*, **275(1)**: 122-124 doi: <https://doi.org/10.1006/abio.1999.4297> (1999).
9. Ehrt, S. and Schnappinger, D., Isolation of plasmids from E. coli by boiling lysis. In *E. coli Plasmid Vectors*. Humana Press, 79-82 doi: <https://doi.org/10.1385/1-59259-409-3:79> (2003).
10. Endres, H. N., Johnson, J. A., Ross, C. A., Welp, J. K. and Etzel, M. R., Evaluation of an ion-exchange membrane for the purification of plasmid DNA. *Biotech and appl biochem*, **37(3)**: 259-266 doi: <https://doi.org/10.1042/BA20030025> (2003).
11. Eon-Duval, A. and Burke, G., Purification of pharmaceutical-grade plasmid DNA by anion-exchange chromatography in an RNase-free process. *Jour of chromat B*, **804(2)**: 327-335 doi: <https://doi.org/10.1016/j.jchromb.2004.01.033> (2004).

12. Eon-Duval, A., MacDuff, R. H., Fisher, C. A., Harris, M. J. and Brook, C., Removal of RNA impurities by tangential flow filtration in an RNase-free plasmid DNA purification process. *Analyt biochem*, **316(1)**: 66-73 doi: [https://doi.org/10.1016/S0003-2697\(03\)00050-2](https://doi.org/10.1016/S0003-2697(03)00050-2) (2003).
13. Ferreira, G. N., Monteiro, G. A., Prazeres, D. M. and Cabral, J. M., Downstream processing of plasmid DNA for gene therapy and DNA vaccine applications. *Trends in biotech*, **18(9)**: 380-388 doi: [https://doi.org/10.1016/S0167-7799\(00\)01475-X](https://doi.org/10.1016/S0167-7799(00)01475-X) (2000).
14. Garger, S. J., Griffith, O. M. and Grill, L. K., Rapid purification of plasmid DNA by a single centrifugation in a two-step cesium chloride-ethidium bromide gradient. *Biochem and biophys res comm*, **117(3)**: 835-842 doi: [https://doi.org/10.1016/0006-291X\(83\)91672-8](https://doi.org/10.1016/0006-291X(83)91672-8) (1983).
15. Giovannini, R. and Freitag, R., Continuous isolation of plasmid DNA by annular chromatography. *Biotech and Bioeng*, **77(4)**: 445-454 doi: <https://doi.org/10.1002/bit.10149> (2002).
16. Guerrero-Germán, P., Prazeres, D. M., Guzmán, R., Montesinos-Cisneros, R. M. and Tejada-Mansir, A., Purification of plasmid DNA using tangential flow filtration and tandem anion-exchange membrane chromatography. *Bioproc and biosys eng*, **32(5)**: 615-623 doi: <https://doi.org/10.1007/s00449-008-0284-7> (2009).
17. Hanora, A., Savina, I., Plieva, F. M., Izumrudov, V. A., Mattiasson, B. and Galaev, I. Y., Direct capture of plasmid DNA from non-clarified bacterial lysate using polycation-grafted monoliths *Jour of biotech*, **123(3)**: 343-355. doi: <https://doi.org/10.1016/j.jbiotec.2005.11.017> (2006).
18. Holmes, D. S. and Quigley, M., A rapid boiling method for the preparation of bacterial plasmids. *Analyt biochem*. **114(1)**: 193-197 doi: [https://doi.org/10.1016/0003-2697\(81\)90473-5](https://doi.org/10.1016/0003-2697(81)90473-5) (1981).
19. Horn, N. A., Meek, J. A., Budahazi, G. and Marquet, M., Cancer gene therapy using plasmid DNA: purification of DNA for human clinical trials. *Hum Gene Ther*, **6(5)**: 565-573 doi: <https://doi.org/10.1089/hum.1995.6.5-565> (1995).
20. Jungbauer, A. and Hahn, R., Polymethacrylate monoliths for preparative and industrial separation of biomolecular assemblies. *Jour of chromat A*, **1184(1-2)**: 62-79. doi: <https://doi.org/10.1016/j.chroma.2007.12.087> (2008).
21. Lander, R. J., Winters, M. A., Meacle, F. J., Buckland, B. C. and Lee, A. L., Fractional precipitation of plasmid DNA from lysate by CTAB. *Biotech and bioeng*, **79(7)**: 776-784 doi: <https://doi.org/10.1002/bit.10335> (2002).
22. Latulippe, D. R. and Zydney, A. L., Size exclusion chromatography of plasmid DNA isoforms. *Jour of Chromat A*, **1216(35)**: 6295-6302 doi: <https://doi.org/10.1016/j.chroma.2009.07.009> (2009).
23. Li, J. F., Li, L. and Sheen, J., Protocol: a rapid and economical procedure for purification of plasmid or plant DNA with diverse applications in plant biology. *Plant Methods*, **6(1)**: 1 doi: <https://doi.org/10.1186/1746-4811-6-1> (2010).
24. Manzano, I., Guerrero-German, P., Montesinos-Cisneros, R. M. and Tejada-Mansir, A., Plasmid DNA pre-purification by tangential flow filtration. *Biotech & Biotechno Equip*, **29(3)**: 586-591 doi: <https://doi.org/10.1080/13102818.2015.1014421> (2015).
25. Murphy, J. C., Wibbenmeyer, J. A., Fox, G. E. and Willson, R. C., Purification of plasmid DNA using selective precipitation by compaction agents. *Nat Biotech*, **17(8)**: 822 doi: <https://doi.org/10.1038/11777> (1999).

26. Sambrook, J. and Russel, D. W., Molecular cloning, a laboratory manual (Third edition), Cold Spring Harbor Laboratory Press (2001).
27. Sandberg, L. M., Bjurling, Å., Busson, P., Vasi, J. and Lemmens, R., Thiophilic interaction chromatography for supercoiled plasmid DNA purification. *Jour of biotech*, **109(1-2)**: 193-199 doi: <https://doi.org/10.1016/j.jbiotec.2003.10.036> (2004).
28. Sasagawa, N., Koebis, M., Yonemura, Y., Mitsuhashi, H. and Ishiura, S., A high-salinity solution with calcium chloride enables RNase-free, easy plasmid isolation within 55 minutes. *Biosci trends*, **7(6)**: 270-275 doi: <https://doi.org/10.5582/bst.2013.v7.6.270> (2013).
29. Sauer, M. L., Kollars, B., Geraets, R. and Sutton, F., Sequential CaCl₂, polyethylene glycol precipitation for RNase-free plasmid DNA isolation. *Analy biochem*, **380(2)**: 310-314 doi: <https://doi.org/10.1016/j.ab.2008.05.044> (2008).
30. Stahl, U., Lemke, P. A., Tudzynski, P., Kück, U. and Esser, K., Evidence for plasmid like DNA in a filamentous fungus, the ascomycete *Podospora anserina*. *Molec and Gener Genet MGG*, **162(3)**: 341-343 doi: <https://doi.org/10.1007/BF00268860> (1978).
31. Sun, B., Yu, X., Yin, Y., Liu, X., Wu, Y., Chen, Y. and Kong, W., Large-scale purification of pharmaceutical-grade plasmid DNA using tangential flow filtration and multi-step chromatography. *Jour of biosc and bioeng*, **116(3)**: 281-286 doi: <https://doi.org/10.1016/j.jbiosc.2013.03.015> (2013).
32. Wahlund, P. O., Gustavsson, P. E., Izumrudov, V. A., Larsson, P. O. and Galaev, I. Y., Polyelectrolyte complexes as a tool for purification of plasmid DNA: Background and development. *Jour of Chromato B*, **807(1)**: 121-127 doi: <https://doi.org/10.1016/j.jchromb.2004.01.046> (2004).